CGCGTTTGGTTGCTCGCTCCACCCCGGAGACCTGGTGTGGTGGAGAAATTTGAA CCCGCAGCCTTAGCTCCGAAAAGGCCGAGTTACCTGGCTCTCCCTGAGTGTCGAG GAGGACATGAGTGAAATGACCAGCGAACTCATTTTTATAGGACTCGGTGAAGC CGGATTCTGCATTTCCCTACTTGTAGACTCATTTTGTGGAATAGAGTTGATCGCTG 5 TCTCCTCCGCAAAGCATTTTAACTCGAATAAGCAAATGCCGCCTCTGTTTGAACG TTTTGGTATTTACAAGAGAAATCATTTTACCTAAGAGAACTAATTGAATTGGC CTGCGAAATAGCAGACGGAGAAATTCCTTTGGAAGTTATTCCGTAGCATAAGAG CTGAAACTTCAGAGCAAGTTTTCATTGGGCAAAATGGGGGAACAACCTATCTTCA 10 GCACTCGAGCTCATGTCTTCCAAATTGACCCAAACACAAAGAAGAACTGGGTAC CCACCAGCAAGCATGCAGTTACTGTGTCTTATTTCTATGACAGCACAAGAAATGT CCCAAACATGACATTTACTAAAACATCTCAGAAGTTTGGCCAGTGGGCTGATAGC CGGGCAAACACCGTTTATGGATTGGGATTCTCCTCTGAGCATCATCTTTCGAAAT 15 TTGCAGAAAAGTTTCAGGAATTTAAAGAAGCTGCTCGACTAGCAAAGGAAAAAT CACAAGAGAAGATGGAACTTACCAGTACACCTTCACAGGAATCCGCAGGCGGG ATCTTCAGTCTCCTTTAACACCGGAAAGTATCAACGGGACAGATGATGAAAGAA CACCTGATGTGACACAGAACTCAGAGCCAAGGGCTGAACCAACTCAGAATGCAT TGCCATTTTCACATAGTTCAGCAATCAGCAAACATTGGGAGGCTGAACTGGCTAC 20 CCTCAAAGGAAATAATGCCAAACTCACTGCAGCCCTGCTGGAGTCCACTGCCAAT GTGAAACAATGGAAACAGCAACTTGCTGCCTATCAAGAGGAAGCAGAACGTCTG CACAAGCGGGTGACTGAACTTGAATGTGTTAGTAGCCAAGCAAATGCAGTACAT ACTCATAAGACAGAATTAAATCAGACAATACAAGAACTGGAAGAGACACTGAAA CTGAAGGAAGAGGAAATAGAAAGGTTAAAACAAGAAATTGATAATGCCAGAGA 25 ACTACAAGAACAGAGGGATTCTTTGACTCAGAAACTACAGGAAGTAGAAATTCG GAACAAGACCTGGAGGGACAACTGTCTGACTTAGAGCAACGTCTGGAGAAAAG TCAGAATGAACAAGAAGCTTTTCGCAATAACCTGAAGACACTCTTAGAAATTCTG GATGGAAAGATATTTGAACTAACAGAATTACGAGATAACTTGGCCAAGCTACTA 30 NNNNNNTTGAATATCACTCCTCCAGGAGGAGGATCTTTTGAAATTGGAATTGTA TATTTCACTGTAAATTTTAGAATCCAGCTTGTAGCTAGTTGGGGAAAAAAGATGA AAAACTTGAACTACAAATTACCTCCATGTATATTATTGGCCATAGTTAACTAGAA AGTTATAAATAGACACTTAATGCAATCTTTTTTCCTGATATTAGCCAATGGGAGA 35 ATTAACAATGTCTAGGTCACATCCCCTTTTTGTGTTCAACACAGTGAAGATTATCT GCTTTTTAAATTAATTTACGATATCTAGAGCTGTGTTTTTGTGCAAAAACTTA GTGATGAAAGCCTGTCTTTTGTTGTAATCTGAATAATTTCTCAGGATATTTTTGCA TGTATCTTTAATTGAAATATACTATAACTGGGTGTATAGAGTTCTTCCCTTTTTTG 40 TGCTGGAAGATATTTCACTCTGGTGACTACTCTGGTACACTCTGGTGTTCTCTAAT CTTGTCTGTTGTATAGTTTACTTTTCCATATTGATTCCATGTATTTATGAGAAGAT ATTGTCTCCCATTTTATTACACATTTTAAAGCCAACTAACGAAGGCAGCTGAGTC CAGCAGCCGTCATTGACAGGCTGGTCTAGCAATGTTAAGTATATTTACAGAATA 45 TGCAGTTACATTTATTATATATTTTGCAAGAAATCTTTTCTGAATGATCAATGCA TTTCAATTACGAATAATAATGGTTATTGGGGAACTGTTTATTATAGATAATTTTA AGGTGTATAGCTATTTTAAAGGGGGTCCATTTACATCAAACAGCCGATCAGAGG ACTCTATCTAAATTGTGATCGTGGCAGATAGAGATGGAGTCATGTACTCTATCTG GCTCTACACATCAATCACATCTTGATTCAAACCTCACAAGGCAATATTCTGAATT

GTTAACTAGGTATTTCAAAACAGGAATTAAATTCAATAGGCTCTTCTCAGTGAAC AGGTTTTAATGTTGTTTTGATGTAATTTTAAAAGACTTTTAGCAAACATGCATTTC TTTATATGATATTTCTTTTACGAAGCTATTTTAAAAAGTAAGCCAAGTGCTGTCT AGTCTGCTTATAAAGTAGGAATTGCATCAGAGTACATATATTCTTGCTGTACAAT 5 GCCTGTGATGTTGAGGAGGGTTCTTTTTAAAGTGTATGCTTGAGTAACTGACTCT ATGGAGTCTATAAATGCACTGACTTCTTGTTTGTACCCCAAAATGATCGAATTGT TAAGTACAAAATTAAGCTAATTAACCAATTTGTAACCATTTTTTCACTCATAAAC 10 CTTAAGTGAGTTTTCAGGTGTCTCTGAAAAATTTATAACAATCATGTATTATATGT GCTGTAACATCATGTACGTTACCTCCATCTATTTTAGGATATTTTCCTCACCTATA TATTATAGGGAGAATAATTTAGATACACATGCTCAGAGCTGAGATATTTCTCTGA TAAATCAGGTAACAAAATGTATTTGATTGATGGAATTTTGAAGTAAATGTGTTTT TATCCATCAGTTTCTGAGTAACAAAGAGCACCAAGTTTTAATTTAAATAGGAGAT 15 TTAACACTAGGGATCAGGGAGTTTAGTATGAAGAGTTAAAAAAATTTAAAAAAC AGTGTAAGCTGTTGAAATGGCAAGTGAATTATTTTAATGATGTAATAAAATATTT TTAAATTTTGACATAGTGATCATTTAATGAAAAAACTCACCAAAATGTCTCCATT TGAATTGTATTGATAATGTGGGACATATGTGTGATTCAATATACATATACCCA TATGTATATACAGAAAATTATTTTTAATACTTTCCTACTGATAATGAAATTTAAAA 20 TTGGAAATTTTGTGAGTGTTTTTCTTGTCCAATAGAGCCTAATTGTTTCCTTTTTTA GTGATTTAACAATCTCTTGAGGGCTGCACCTTTAAATTCCCAGATTGTCAATAGA CTTAAGACTTTTAACTATTCATTTACAGTAGGAGAGTATGTAGAAATCATCATCC ACAGTCATAATTAGGTTGTGTGCCTACTGTAGTTTTTCCATTTCTGTATTATAT 25 AAACATTTGCATATTAAAATTTGATTTTTCCCAGAGACAAGTATTATATACTGTAT CTATATTTAAATCAAACTGTGGTAATATATTTCTCAGAAAATAATGTTGGGGACT ATAGCCTGAACATGTGGACTTGAAGCGACATGGAGGAGGAGGTTGATCCCATTG TGTATAAGTTAATATGTGATAACTATTGAATCTTGTACAAAAAACAAAAATTGANA 30 AGAATGTACTTTATCTTTTTTCCTCCAGTCTTTTACAGATATTTAAAAGCATTTA AATGATGACAGCATTTACTTAAATCTTTCAGGTGCTACTGGATTTTGCATTAGTGT GTTATGTTGTGAAATCCTAACTTTGACATAAAAGGTTTTATAAGTATTCCCCTGCC TGGAAAATTAGTTTTTCTCCNCTCTCTCTCTTTTCTCTTTTTCTG CAGACTAAAACATGCTCACGAAGTTGCATCTCTCTCTTGTCTCTATAGAAGATCTC 35 CAGCACCATCATAGATTTGATGTTCTGCTGTCATTGNACTGTTGGGAAGCAGTTA GAGGAAAAGCTCACTTTTTTTTCAGGTGGAAATAAAAGGAACACTCAAAATTA AGCCAACACCACCACTTTAAAAACTAGTTTATTTGCCCTGTTAAAATTAAA TGATTCTTNAACATGTGGGCTACAGTCTCCCATGTTTTTATTTAACTGAAGCATAT ACACTTCGGNCATTTATCTCCTGTGGNCCTGATTTTGTCAGTACTGGAATG

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SEQ ID NO: 654
>21590 BLOOD INCYTE\_3985758H1
GCNACGGTTGGCGCTCGNCCTGGAGCCTGCCCTGGCGTNCCCCCGCGGGCGCAG
CCAAGCTTCTTGGCNATGGTAGATAACTGCAGGGGACTCTGGCCGCGGCTAACTA
NCCTGGAGATGCTGATCGGGACCCCCCCGCAGAAGCTACAGATTCTCGTTGACA
NTGGAAGCAGTAACTTTGA

SEQ ID NO: 655

>21591 BLOOD 404604.3 AF122922 g4585369 Human Wnt inhibitory factor-1 mRNA, complete cds. 0

- 15 CTGAAGGCAACACCATTCTCCAAACACCTCAAAATGCTATCTTCTTTAAAACATG
  TCAACAAGCTGAGTGCCCAGGCGGGTGCCGAAATGGAGGCTTTTGTAATGAAAG
  ACGCATCTGCGAGTGTCCTGATGGGTTCCACGGACCTCACTGTGAGAAAGCCCTT
  TGTACCCCACGATGTATGAATGGTGGACTTTGTGTGACTCCTGGTTTCTGCATCTG
  CCCACCTGGATTCTATGGAGTGAACTGTGACAAAGCAAACTGCTCAACCACCTGC
- 20 TTTAATGGAGGACCTGTTTCTACCCTGGAAAATGTATTTGCCCTCCAGGACTAG AGGGAGAGCAGTGTGAAATCAGCAAATGCCCACAACCCTGTCGAAATGGAGGTA \*AATGCATTGGTAAAAGCAAATGTAAGTGTTCCAAAGGTTACCAGGGAGACCTCT !GTTCAAAGCCTGTCTGCGAGCCTGGCTGTGGCACATGGAACCTGCAATAAAAGGTA
- 25 CGAAGCCAGCCTCATACATGCCCTGAGGCCAGCAGGCGCGCCAGCTCAGGCAGCA CACGCCTTCACTTAAAAAGGCCGAGGAGCGGCGGGATCCACCTGAATCCAATTA CATCTGGTGAACTCCGACATCTGAAACGTTTTAAGTTACACCAAGTTCATAGCCT TTGTTAACCTTTCATGTGTTGAATGTTCAAATAATGTTCATTACACTTAAGAATAC TGGCCTGAATTTTATTAGCTTCATTATAAATCACTGAGCTGATATTTACTCTTCCT
- TTTAAGTTTTCTAAGTACGTCTGTAGCATGATGGTATAGATTTTCTTGTTTCAGTG CTTTGGGACAGATTTTATATTATGTCAATTGATCAGGTTAAAAATTTTCAGTGTGTA GTTGGCAGATATTTTCAAAATTACAATGCATTTATGGTGTCTGGGGGCAGGGGAA CATCAGAAAGGTTAAATTGGGCAAAAATGCGTAAGTCACAAGAATTTGGATGGT GCAGTTAATGTTGAAGTTACAGCATTTCAGATTTTATTGTCAGATATTTAGATGTT
- 40 GCTTTAGTTTCTGAGCATTGTGTGGAGGTNANCTTTGCACATGCTATCTTATGAA AATAAAATTGGTTGCAATTTAGTGGT

**SEQ ID NO: 656** 

- >21600 BLOOD 480735.6 U60477 g1575342 Human apolipoprotein AI regulatory protein-1/chicken ovalbumin upstream promoter transcription factor II (TFCOUP2) gene, complete cds. 0
  - CATCGAGTGCGTGTGTGCGGAGACAAGTCGAGCGCAAGCACTACGGCCAGTT CACGTGCGAGGGCTGCAAGAGCTTCTTCAAGCGCAGCGTGCGGAGGAACCTGAG CTACACGTGCCGCCCAACCGGAACTGTCCCATCGACCAGCACCATCGCAACCA

SEQ ID NO: 658 >21621 BLOOD 253228.8 Incyte Unique

- 30 GAGCCTGCTGGGGACCCAGGTATTCTTCCTTCTGGGGACCCTGGGCCTCTTCTGC
  CTCGTGTTTGCCTGTGTGAAGCCCGACTTCTCCACCTGTGCCTCTCGGCGCTT
  CCTCTTTGGGGTTCTGTTCGCCATCTGCTTCTTGTCTGGCGGCTCACGTCTTTGC
  CCTCAACTTCCTGGCCCGGAAGAACCACGGGCCCCGGGGCTGGTGATCTTCACT
  GTGGCTCTGCTGCTGACCCTGGTAGAGGTCATCAATACAGAGTGGCTGATCA
  35 TCACCCTGGTTCGGGGCAGTGGCGAGGGCGCCCTCAGGGCAACAGCAGCGCAG
- 40 GCAGCACAACAGTCCCACCTGGGATGACCCCACGCTGGCCATCGCCCTCGCCGCC
  AATGCCTGGGCCTTCGTCCTCTTCTACGTCATCCCCGAGGTCTCCCAGGTGACCA
  AGTCCAGCCCAGAGCAAAGCTACCAGGGGGACATGTACCCCACCCGGGGCGTGG
  GCTATGAGACCATCCTGAAAGAGCAGAAGGGTCAGAGCATGTCGTGGAGAACA
  AGGCCTTTTCCATGGATGAGCCGGTTGCAGCTAAGAGGCCGACTCAGATGCCCT
- 45 CGGGTACAATGGGCAGCTGCTGACCAGTGTGTACCAGCCCACTGAGATGGCCCT GATGCACAAAGTTCCGTCCGAAGGAGCTTACGACATCATCCTCCCACGGGCCACC GCCAACAGCCAGGTGATGGGCAGTGCCAACTCGACCCTGCGGGCTGAAGACATG TACTCGGCCCAGAGCCACCAGGCGGCCACACCGCCGAAAGACGGCAAGAACTCT CAGGTCTTTAGAAACCCCTACGTGTGGGACTGAGTCAGCGGTGGCGAGGAGAGG

5

10 **SEO ID NO: 659** >21628 BLOOD 255990.10 AJ011497 g4128014 Human mRNA for Claudin-7. 0 GCCGGAGGGGACAGTGGTAGGTGGGGAGGTTGAGTGCAAAGGGTTCAGGCTGTA AGTCATGTTGGGTTGGAATGGGGGCACAGGAAGGTGGGGCTGTTGGGGAGCCAC GCTAAGCCGGGTGTCTGTAGCAGAGCCAGAGAACCGGGACACTGAAGAGGGTGC 15 TGAAGGGGGCGACTCTCAGGGATCGAGCCAGGGCCCCCGAAGGTGGGATCGACC AGGGTAGGAGACAGGAAAAAAAAGGAGAGCAGCGGGTGGGGGCGAAAGCAGG GCCGAGGAGAGACACTTTGGACAGAACCCGGCGGGAAAGGGCGCGCCGAG GCTTGTCAGGGGCGCCCCGCAGCGTCCCAGGCGCACCTGTTGGGAAGAAAGGAA GGGGCTTCCCGGTGTTCGAGGGAAATCCAGTCCGGAGGGGCTGACTCGGAGCTT 20 GGGACTCCTGGGGAGCCACCGCCTCCTCCCCAGCGGCGGTCAAAACCGGGCAAG CGAAGGGGCGTGACCCTGGTGCTCAGGTTTCTTCCTCCTCACCTGGGCAAGGAGG GGTGGGGCCACGACTTCCGGTTCAGGTGAGTGTCCCTTCGGTGACGTCAGGTCA ATCCTCGGCCGCCCTCCGGTCCCGCCTCCCCTCCCGCGCTCCCGGGGCGCGCGG 25 CCTGCTGGCTCACCTCCGAGCCACCTCTGCTGCGCACCGCAGCCTCGGACCTACA GCCCAGGATACTTTGGGACTTGCCGGCGCTCAGAAACGCGCCCAGACGGCCCCT CCACCTTTTGTTTGCCTAGGGTCGCCGAGAGCGCCCGGAGGGAACCGCCTGGCCT TCGGGGACCACCATTTTGTCTGGAACCACCCTCCGGCGTATCCTACTCCCTGT GCCGCGAGGCCATCGCTTCACTGGAGGGGTCGATTTGTGTGTAGTTTGGTGACAA 30 GATTTGCATTCACCTGGCCCAAACCCTTTTTGTCTCTTTTGGGTGACCGGAAAACTC GGTCTCCCGCCGGCGCCCCAGTGTTTTCTGAGGGCGGAAATGGCCAATTCGG CCTGCACCGCCATCCCGCAGTGGCAGATGAGCTCCTATGCGGGTGACAACATCAT 35 CACGCCCAGGCCATGTACAAGGGGCTGTGGATGGACTGCGTCACGCAGAGCAC GGGGATGATGAGCTGCAAAATGTACGACTCGGTGCTCGCCCTGTCCGCGGCCTTG CAGGCCACTCGAGCCCTAATGGTGGTCTCCCTGGTGCTGGGCTTCCTGGCCATGT TTGTGGCCACGATGGGCATGAAGTGCACGCGCTGTGGGGGAGACGACAAAGTGA AGAAGGCCCGTATAGCCATGGGTGGAGGCATAATTTTCATCGTGGCAGGTCTTGC 40 CGCCTTGGTAGCTTCCTGGTATGGCCATCAGATTGTCACAGACTTTTATAACC CTTTGATCCCTACCAACATTAAGTATGAGTTTGGCCCTGCCATCTTTATTGGCTGG GCAGGGTCTGCCCTAGTCATCCTGGGAGGTGCACTGCTCTCTGTTCCTGTCCTG GGAATGAGAGCAAGGCTGGGTACCGTGTACCCCGCTCTTACCCTAAGTCCAACTC TTCCAAGGAGTATGTGTGACCTGGGATCTCCTTGCCCCAGCCTGACAGGCTATGG 45 GAGTGTCTAGATGCCTGAAAGGGCCTGGGGCTGAGCTCAGCCTGTGGGCAGGGT GCCGGACAAAGGCCTCCTGTCACTCTGTCCCTGCACTCCATGTATAGTCCTCTT GGGTTGGGGGGGGGGTGCCGTTGGTGGAGAGACAAAAAGAGGGAGAGTG TGCTTTTTGTACAGTAATAAAAAATAAGTATTGGGAAGCAGGCTTTTTTCCCTTC AGGGCCTCTGCTTCCTCCCGTCCAGATCCTTGCAGGGAGCTTGGAACCTTAGTG

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SEQ ID NO: 660 >21631 BLOOD 370788.1 AK000072 g7019922 Human cDNA FLJ20065 fis, clone COL01613, highly similar to ECLC\_BOVIN EPITHELIAL CHLORIDE CHANNEL PROTEIN. 0

- - 25 CAAAACATAAAGTGCAATTTTAGAAGTACATGGGAGGTGATTAGCAATTCTGAG GATTTTAAAAACACCATACCCATGGTGACACCACCTCCTCCACCTGTCTTCTCATT GCTGAAGATCAGTCAAAGAATTGTGTGCTTAGTTCTTGATAAGTCTGGAAGCATG GGGGGTAAGGACCGCCTAAATCGAATGAATCAAGCAGCAAAACATTTCCTGCTG CAGACTGTTGAAAAATGGATCCTGGGTGGGGATGGTTCACTTTGATAGTACTGCCA

  - 35 AGTAATAGAGATGAGCAAGATAACAGGAGGAAGTCATTTTTATGTTTCAGATGA AGCTCAGAACAATGGCCTCATTGATGCTTTTGGGGCTCTTACATCAGGAAATACT GATCTCTCCCAGAAGTCCCTTCAGCTCGAAAGTAAGGGATTAACACTGAATAGTA ATGCCTGGATGAACGACACTGTCATAATTGATAGTACAGTGGGAAAGGACACGT TCTTTCTCATCACATGGAACAGTCTGCCTCCCAGTATTTCTCTCTGGGATCCCAGT
  - 40 GGAACAATAATGGAAAATTTCACAGTGGATGCAACTTCCAAAATGGCCTATCTC AGTATTCCAGGAACTGCAAAGGTGGGCACTTGGGCATACAATCTTCAAGCCAAA GCGAACCCAGAAACATTAACTATTACAGTAACTTCTCGAGCAGCAAATTCTTCTG TGCCTCCAATCACAGTGAATGCTAAAAATGAATAAGGACGTAAACAGTTTCCCCA GCCCAATGATTGTTTACGCAGAAAATTCTACAAGGATATGTACCTGTTCTTGGAGC
  - 45 CAATGTGACTGCTTTCATTGAATCACAGAATGGACATACAGAAGTTTTGGAACTT
    TTGGATAATGGTGCAGGCGCTGATTCTTTCAAGAATGATGGAGTCTACTCCAGGT
    ATTTTACAGCATATACAGAAAATGGCAGATATAGCTTAAAAGTTCGGGCTCATGG
    AGGAGCAAACACTGCCAGGCTAAAATTACGGCCTCCACTGAATAGAGCCGCGTA
    CATACCAGGCTGGTAGTGAACGGGGAAATTGAAGCAAACCCGCCAAGACCTGA

AATTGATGAGGATACTCAGACCACCTTGGAGGATTTCAGCCGAACAGCATCCGG AGGTGCATTTGTGGTATCACAAGTCCCAAGCCTTCCCTTGCCTGACCAATACCCA CCAAGTCAAATCACAGACCTTGATGCCACAGTTCATGAGGATAAGATTATTCTTA CATGGACAGCACCAGGAGATAATTTTGATGTTGGAAAAGTTCAACGTTATATCAT 5 AAGAATAAGTGCAAGTATTCTTGATCTAAGAGACAGTTTTGATGATGCTCTTCAA GTAAATACTACTGATCTGTCACCAAAGGAGGCCAACTCCAAGGAAAGCTTTGCA AAAGTATAGATAAAAGCAATTTGACATCAAAAGTATCCAACATTGCACAAGTAA CTTTGTTTATCCCTCAAGCAAATCCTGATGACATTGATCCTACACCTACTTCCTAC 10 TCCTACTCCTACTCCTGATAAAAGTCATAATTCTGGAGTTAATATTTCTACGCTGG TATTGTCTGTGATTGGGTCTGTTGTAATTGTTAACTTTATTTTAAGTACCACCATT TGAACCTTAACGAAGAAAAAATCTTCAAGTAGACCTAGAAGAGAGTTTTAAAA AACAAAACAATGTAAGTAAAGGATATTTCTGAATCTTAAAATTCATCCCATGTGT GATCATAAACTCATAAAAATAATTTTAAGATGTCGGAAAAGGATACTTTGATTAA 15 ATAAAAACACTCATGGATATGTAAAAACTGTCAAGATTAAAATTTAATAGTTTCA TTTATTTGTTATTTGTAAGAAATAGTGATGAACAAGATCCTTTTTCATAC TGATACCTGGTTGTATATTTGATGCAACAGTTTTCTGAAATGATATTTCAAAT TGCATCAAGAAATTAAAATCATCTATCTGAGTAGTCAAAATACAAGTAAAGGAG AGCAAATAAACAACATTTGGAAAAAAATG

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SEO ID NO: 661

A 1/2013 >24.656 BLOOD INCYTE 547531H1 (1901) A 1/2014 (1901) A 1/2014 (1901) A 1/2014 (1901) A 1/2014 (1901) THE ACCIDATE STREET OF THE STR \*\*\*\*\*\*\*\*AGGANTTGTTCAGGGGTTAAGAAAGGGTTTGAGGTGGTTGAAAAACTGAATGGA

AAAGCTTATGGCTCTGTGATGATATTAGTGACCAGCGGAGATGATAAGCTTCTTG 25 GCAATTGCTTACCCACTGTGCTCAGCAGTGGTTCAACAATTCACTCCATTGCCCT GGGTTCATCTGCAGCCCCAAATCTGGA

**SEO ID NO: 662** 

- 30 >21660 BLOOD 238908.1 AL137516 g6808175 Human mRNA; cDNA DKFZp564M2178 (from clone DKFZp564M2178); partial cds. 0 GAACCACCGGCAGACGCACCTCCGGGCCACACCCACGAGGCTCCTGCCCCTGTT GTCCTGGGGTCCCCAGTTGTTCTAGGGCCTCCTGTGGGCCAGGCCCGAGTGGCTG TGGAGCACTCATACCGAAAGGCAGAAGAGGGTGGGGAAGGGGCGACTGTCCCAT 35 CTGCCGCTGCCACCACCACTGAGGTAGTGACTGAGGTGGAGCTGCTCCTCTACAA GTGCTCTGAGTGCTCCCAGCTCTTCCAGCTGCCGGCGGATTTCCTGGAGCACCAG AGGTGCAGGCCTCGTCACCTGCAGAGGTGCCTGTGTCTCAGCCTGACCCCTTGCC AGCTTCTGACCACAGTTACGAGCTGCGCAATGGTGAAGCCATTGGGCGGGATCG 40 CCGGGGGCCAGGGCCCGGAGGAACAACAGTGGAGAAGCAGGCGGGCAGCCA CACAGGAGCTCTTCTGCTCAGCCTGTGACCAGCTCTTTCTCTCACCCCACCAGCTA CAGCAGCACCTGCGGAGTCACCGGGAGGGCGTCTTTAAGTGCCCCCTGTGCAGTC GTGTCTTCCCTAGCCCTTCCAGTCTGGACCAGCACCTTGGAGACCATAGCAGCGA
- GTCACACTTCCTGTGTGTAGACTGTGGCCTGGCCTTCGGCACAGAGGCCCTCCTC 45 CTGGCCCACCGGCGAGCCCACACCCCGAATCCTCTGCATTCATGTCCATGTGGGA AGACCTTTGTCAACCTTACCAAGTTCCTTTATCACCGGCGTACTCATGGGGTAGG GGGTGTCCCTCTGCCCACAACACCAGTCCCACCAGAGGAACCTGTCATTGGTTTC CCTGAGCCAGCCCAGCAGAGACTGGAGAGCCAGAGGCCCCTGTG TCTGAGGAGACCTCAGCAGGGCCCGCTGCCCCAGGCACCTACCGCTGCCTCTGT

GCAGCCGTGAATTTGGAAAGGCCTTGCAGCTGACCCGGCACCAACGTTTTGTGCA TCGGCTGGAGCGCCCATAAATGCAGCATTTGTGGCAAGATGTTCAAGAAGAA GTCTCACGTGCGTAACCACCTGCGCACACACACGGGGAGCGGCCCTTCCCCTGC CCTGACTGCTCCAAGCCCTTCAACTCACCTGCCAACCTGGCCCGCCACCGGCTCA 5 CACACAGGAGAGGGCCCTACCGGTGTGGGGACTGTGGCAAGGCTTTCACGC AAAGCTCCACACTGAGGCAGCACCGCTTGGTGCATGCCCAGCACTTCCCCTACCG  ${\tt CTGCCAGGAATGTGGGGTGCGTTTTCACCGTCCTTACCGCCTGCTCATGCACCGC}$ TACCATCACAGGTGAATACCCCTACAAGTGTCGCGAGTGCCCCCGCTCCTTCT TGCTGCGTCGGCTGCAGGTGCACCAGCTCGTGGTCCATGCCGGGCGCCAGCC 10 CCACCGCTGCCCATCCTGTGGGGCTGCCTTCCCCTCCTCACTGCGGCTCCGGGAG CACCGCTGTGCAGCCGCTGCTGCCCAGGCCCCACGGCGCTTTGAGTGTGGCACCT GTGGCAAGAAAGTGGGCTCAGCTGCTCGACTGCAGGCACACGAGGCGCCCATG CAGCTGCTGGGCCTGGAGAGGTCCTGGCTAAGGAGCCCCCTGCCCCTCGAGCCCC ACGGGCCACTCGTGCACCAGTTGCCTCTCCAGCAGCCCTTGGAAGCACTGCTACA 15 GCATCCCCTGCGGCCCCGCCGCCGGGGTCTAGAGTGCAGCGAGTGCAAG AAGCTGTTCAGCACAGAGACGTCACTGCAGGTGCACCGGCGCATCCACACAGGT GAGCGCCATACCCATGTCCAGACTGTGGCAAAGCGTTCCGTCAGAGTACCCAC CTGAAAGACCACCGGCGCCTGCACACAGGTGAGCGGCCCTTTGCCTGTGAAGTG TGTGGCAAGGCCTTTGCCATCTCCATGCGCCTGGCAGAACATCGCCGCATCCACA 20 CAGGCGAACGACCCTACTCCTGCCCTGACTGTGGCAAGAGCTACCGCTCCTTCTC CAACCTCTGGAAGCACCGCAAGACCCATCAGCAGCAGCATCAGGCAGCTGTGCG .GCAGCAGCTGGCAGAGGCGGAGGCTGCCGTTGGCCTGGCCGTCATGGAGACTGC TGTGGAGGCGCTACCCCTGGTGGAAGCCATTGAGATCTACCCTCTGGCCGAGGCT · GAGGGGTCCAGATCAGTGGCTGACTCTGCCCGACTTCCTCTTTGGCACCTCCAT 25 TCCCTGTTGCTGAAGGCCCTCCAGCATCCCCTTAAGCATCTGTACATACTGTGTCC CTTCCTCTTCCCATCCCACCACCTTGTAAGTTCTAAATTGGATTTATTCTCTCGT CTCTTAGCACTGGTGACCCCAAAAATGAAACCATCAATAAAGACTGAGTTGCC

30 **SEQ ID NO: 663** >21669 BLOOD 132774.1 Incyte Unique GCCGGACAGAGCAGAAGCCCTCTTGGACTGGACGATTTGGGAATTCAAAACT TGGGACAAACTGTCAGCCTTGCCCCTGCTGTGGAGGCAGCCTCAATGCTGAAAAT GGAGCCTCTGAACAGCACGCACCCGGCACCGCCTCCAGCAGCCCCCTGGA 35 GTCCCGTGCGGCGGTGGCGGCAGCGCAATGGCAACGAGTACTTCTACATTCTG GTTGTCATGTCCTTCTACGGCATTTTCTTGATCGGAATCATGCTGGGCTACATGAA ATCCAAGAGGCGGGAGAAGAAGTCCAGCCTCCTGCTGCTGTACAAAGACGAGGA GCGGCTCTGGGGGGAGGCCATGAAGCCGCTGCCCGTGGTGTCGGGCCTGAGGTC GGTGCAGGTGCCCCTGATGCTGAACATGCTGCAGGAGAGCGTGGCGCCCGCGCT 40 GTCCTGCACCCTCTGTTCCATGGAAGGGGACAGCGTGAGCTCCGAGTCCTCCC CCGGACGTGCACCTCACCATTCAGGAGGAGGGGGGCAGACGAGGAGCTGGAGGA GACCTCGGAGACGCCCTCAACGAGAGCAGCGAAGGGTCCTCGGAGAACATCCA CTTAGAGAGAGAAAGACAGTTTTCAAGTGTCTGGTTTCACTTTCACAGTGCGGC 45 AGGCTCAGCCGGAACCAGCACCTCCAAGGAGTCCGGGAGGTGCCTGTGGTTTAC ACCCACCACTGAAAAAGCCGCGGAGATGCGCAGCGCGTACACTGACTTTGGGGC CTGGGTGTTGGGGTTCTGATCAGAATTTGGCGGGATGATATGCTTGCCATTTTCTC

ACTGGATGCCCTGGGTAGCTCCTGCAGGGTCTGCCTGTTCCCAGGGCTGCCGAAT

GCTTAGGACACGCTGAGAGACTAGTTGTGATTTGCTATTTTGCCTAGAGCTTTGT CCTTCTAGATCTGATTGGCTGTAAGTATCTCTACTGTGTACCTGTGGCATTCCTTC ACAGTGGGTTACAAGCTTCTTTGGGATTAGAGGGGGATTTTGGATGGGAGAAAG CGTGGGAGATCGTGGAACCCCAGCCCCATTTGCACACTATAAGAAAAAAAGTAA CTTTTAAACCTGTTAACATTGGCCGGGGTTATAAGAGATGATCTTCTATTT

**SEQ ID NO: 664** 

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>21683 BLOOD 444662.14 Z58148 g1029379 Human CpG island DNA genomic Mse1 fragment, clone 30a7, forward read cpg30a7.ft1d. 3e-15

- CTCAGCGTCAGGCAAGTTGGCCTCTCTGTTGTAAATTAGTGGTTAAGGTTATCTA
   TTATTGCCACTTTTCCAGCGCTAAAGGCTGTTTTTGGAACCAGTGTTGCTTGTTCCG
   CGGGTGATTGGCTTTTTTTTTTTTTTGCAAACCAGTTATTCAAGTTTCTGGTCTTTAA
   AAAACTCTGTGGCGGTACGGTAACCGAGGAGGTTCCAGCGCGGGGAAGTACCC
   CGCGGGTGGGTGTGTGCGCAAGGCCAGGGCCAGAGGGCACGTGGCGCCGGGA
   GGAGAGAGAATGTCTTTTCGAGGCGGAGGTCGTGGAGGCTTTAATCGAGGTGGT
   GGAGGTGGCCGGCTTCAACCGAGGCGGCAGCAACCACTTCCGAGGTGGAG
   CGGCGGTGGAGGCGGCAATTTCAGAGGCGGCGCAGGGGAGGATTTGGAC
   GAGGGGGTGGCCGCGGAGGCTTTAACAAAGGCCAAGACCAAGGACCTCCAGAA
   CGTGTAGTCTTATTAGGAGAGTTCCTGCATCCCTGTGAAGATGACATAGTTTGTA
   AATGTACCACAGATGAAAATAAGGTGCCTTATTTCAATGCTCCTGTTTACTTAGA
   AAACAAAGAACAAATTGGAAAAAGTGGATGAAAATATTTGGACAACTCAGAGATTT

**SEQ ID NO: 665** 

SEQ ID NO: 666

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>21694 BLOOD 029567.1 Incyte Unique

GCCACCACCCAGCTGCTTAAGCACAAACTAATTTCAAAACCAGTCTTAGAAAT TATATCCTACGCACTTGTCAAACGGGGTCAGTTTTTCTTGAAAGTAAACCTCTGCT

CTTCATCACACAATCTAAATCTGCCACCCTACCTAAGGCAGGGACTTAAAATGAG GGGCAGGTTTTCTCAGATAAAATAAGCAAACAGACGAATTGGAATATTTCGTCTC AATTCCCATGTACAATTTTCAGCCTCATATGCAAATCAATATGGCAACCATCTCTT TTTTCTATCAGCAAGAGCCATGTGTTGGTATTAAGAGGCTAGGTTGTAGTTCCCC 5 TCTTGACACCAGAAACACCAGGCCTACTGTTTTGTTTGATAGCCTAGCACAGATG TAACTCTTCTAAAGGGTAACATTTACTACTACACAGAAAGTCATTTTTAGAATGT TCCTAGTCCATCCAAGAAAGGCTAAAAAATGTTCTGTGTGATTCTGGACTTAAGA AGTCTTTTCACTGAATTCTGGCCCTAGATGCTCAACTAGTAACCATAATGC CCTGTTTTTCCCAAATGCTGAAATGGAACATTTACTGCATTGGCAATGTTTTCTAG 10 TGGGATTGGTTACAGAAACTGTCATTCATTCTTACTGTGCAATATTACACAGCCA ATACATTGTAAGTAAGAAAATATACAGGTTAGAAAACTGTAGGTATAGCATATT GTAAAAAGAAAATATATATACACGCATAGCGAAAAATGTCAGGAAGAATATACC AAAATGTTAACGATTATCTTTAAGTAGTATCATCTTTTTTACTTCTTGTCTTCTGT 15 TTCTCCTAAAACTTCTAAAATAAAAAGGTATTATTTTCAT

**SEQ ID NO: 667** >21697 BLOOD 350207.6 X69086 g34811 Human mRNA for utrophin. 0 GCGGGCAGCAGCCGCCGCGGGCTTTCTCCCGCCGAGGGGCGAGGAGGAGC 20 CTCTGGCTCCAGAAGCCGATTGGGGAATCACGGGGAGCGGCCCCCCTTCTTT GGGTCATTTCTGCAAACGGAAAACTCTGTAGCGTTTGGCAAAGTTGGTGCCTGCN CGCCCTTCCAGGTTTGCGCTTTGACTGTTTTGTTTTTGGCGGAACTACCAGGCAG 25 TAAAATACATCGCACCACCAAACTAACACTCGCACACCCCCCGCGGTTACTCCG TGTCAAACTCCTAGAGGAGCCCTTGGCCAGCTCGGGGTGCGGCGGTGGCGACCG GCAGGCGAGGAGCCCGCGGGCAGCAGGTATTGATGTCAAGCTGAACCATCGTA GGAAGTTGAAAGCCTTAGAAAGAGGACTTGGTAAAGTTTTTGGATTATCTTGAAA CTCTGGCAAAATGGCCAAGTATGGAGAACATGAAGCCAGTCCTGACAATGGGCA 30 GAACGAATTCAGTGATATCATTAAGTCCAGATCTGATGAACACAATGACGTACA GAAGAAAACCTTTACCAAATGGATAAATGCTCGATTTTCAAAGAGTGGGAAACC CTAGAAGGCCTCACAGGAACATCACTGCCAAAGGAACGTGGTTCCACAAGGGTA CATGCCTTAAATAACGTCAACAGAGTGCTGCAGGTTTTACATCAGAACAATGTGG 35 AATTAGTGAATATAGGGGGAACTGACATTGTGGATGGAAATCACAAACTGACTT TGGGGTTACTTTGGAGCATCATTTTGCACTGGCAGGTGAAAGATGTCATGAAGGA TGTCATGTCGGACCTGCAGCAGACGAACAGTGAGAAGATCCTGCTCAGCTGGGT GCGTCAGACCACCAGGCCTACAGCCAAGTCAACGTCCTCAACTTCACCACCAGC TGGACAGATGGACTCGCCTTTAATGCTGTCCTCCACCGACATAAACCTGATCTCT 40 TCAGCTGGGATAAAGTTGTCAAAATGTCACCAATTGAGAGACTTGAACATGCCTT CAGCAAGGCTCAAACTTATTTGGGAATTGAAAAGCTGTTAGATCCTGAAGATGTT GGTGCTACCTCAGCAAGTCACCATAGACGCCATCCGTGAGGTAGAGACACTCCC AAGGAAATATAAAAAAGAATGTGAAGAAGAGGCAATTAATATACAGAGTACAG 45 CGCCTGAGGAGGAGCATGAGAGTCCCCGAGCTGAAACTCCCAGCACTGTCACTG AGGTCGACATGGATCTGGACAGCTATCAGATTGCGTTGGAGGAAGTGCTGACCT GGTTGCTTTCTGAGGACACTTTCCAGGAGCAGGATGATATTTCTGATGATGT TGAAGAAGTCAAAGACCAGTTTGCAACCCATGAAGCTTTTATGATGGAACTGACT 

CAAGGAACTCTGTCAGACGAAGAAGAATTTGAGATTCAGGAACAGATGACCCTG CTGCACGATGTGCTGATGGAACTGCAGAAGAAGCAACTGCAGCAGCTCTCCGCC TGGTTAACACTCACAGAGGAGCGCATTCAGAAGATGGAAACTTGCCCCCTGGAT 5 GATGATGTAAAATCTCTACAAAAGCTGCTAGAAGAACATAAAAGTTTGCAAAGT GATCTTGAGGCTGAACAGGTGAAAGTAAATTCACTAACTCACATGGTGGTCATTG TTGATGAAAACAGTGGTGAGAGCGCTACAGCTATCCTAGAAGACCAGTTACAGA AACTTGGTGAGCGCTGGACAGCAGTATGCCGTTGGACTGAAGAACGCTGGAATA GGTTACAAGAAATCAATATTGTGGCAGGAATTATTGGAAGAACAGTGCTTGTT 10 CTTCAAAGACCAAAAGGAACTAAGTGTCAGTGTTCGACGTCTGGCTATTTTGAAG GATGTGGGACAATTACTTGATAATTCCAAGGCATCTAAGAAGATCAACAGTGAC TCAGAGGAACTGACTCAAAGATGGGATTCTTTGGTTCAGAGACTAGAAGATTCCT 15 CCAACCAGGTGACTCAGGCTGTAGCAAAGCTGGGGATGTCTCAGATTCCTCAGA AGGACCTTTTGGAGACTGTTCGTGTAAGAGAACAAGCAATTACAAAAAATCTA AGCAGGAACTGCCTCCTCCTCCCCCAAAGAAGAGACAGATCCATGTGGATAT TGAAGCTAAGAAAAGTTTGATGCTATAAGTGCAGAGCTGTTGAACTGGATTTTG AAATGGAAAACTGCCATTCAGACCACAGAGATAAAAGAGTATATGAAGATGCAA 20 GACACTTCCGAAATGAAAAAGAAGTTGAAGGCATTAGAAAAAAGAACAGAGAGA AAGAATCCCCAGAGCAGATGAATTAAACCAAACTGGACAAATCCTTGTGGAGCA AATGGGAAAAGAAGGCCTTCCTACTGAAGAAATAAAAAAATGTTCTGGAGAAGGT GTCATCAAGACAAAGGAGGAGTGGGTAAAACACACTTCCATTTCTGAATCTTCCC 25 GGCAGTCCTTGCCAAGCTTGAAGGATTCCTGTCAGCGGGAATTGACAAATCTTCT TGGCCTTCACCCCAAAATTGAAATGGCTCGTGCAAGCTGCTCGGCCCTGATGTCT CAGCCTTCTGCCCCAGATTTTGTCCAGCGGGGCTTCGATAGCTTTCTGGGCCGCT ACCAAGCTGTACAAGAGGCTGTAGAGGATCGTCAACAACATCTAGAGAATGAAC 30 TGAAGGGCCAACCTGGACATGCATATCTGGAAACATTGAAAACACTGAAAGATG TGCTAAATGATTCAGAAAATAAGGCCCAGGTGTCTCTGAATGTCCTTAATGATCT TGCCAAGGTGGAGAAGGCCCTGCAAGAAAAAAAGACCCTTGATGAAATCCTTGA GAATCAGAAACCTGCATTACATAAACTTGCAGAAGAAACAAAGGCTCTGGAGAA AAATGTTCATCCTGATGTAGAAAAATTATATAAGCAAGAATTTGATGATGTGCAA 35 GGAAAGTGGAACAAGCTAAAGGTCTTGGTTTCCAAAGATCTACATTTGCTTGAGG AAATTGCTCTCACACTCAGAGCTTTTGAGGCCGATTCAACAGTCATTGAGAAGTG GATGGATGGCGTGAAAGACTTCTTAATGAAACAGCAGGCTGCCCAAGGAGACGA CGCAGGTCTACAGAGGCAGTTAGACCAGTGCTCTGCATTTGTTAATGAAATAGAA ACAATTGAATCATCTCTGAAAAACATGAAGGAAATAGAGACTAATCTTCGAAGT 40 GGTCCAGTTGCTGGAATAAAAACTTGGGTGCAGACAAGACTAGGTGACTACCAA ACTCAACTGGAGAAACTTAGCAAGGAGATCGCTACTCAAAAAAGTAGGTTGTCT GAAAGTCAAGAAAAGCTGCGAACCTGAAGAAAGACTTGGCAGAGATGCAGGA ATGGATGACCCAGGCCGAGGAAGAATATTTGGAGCGGGATTTTGAGTACAAGTC ACCAGAAGAGCTTGAGAGTGCTGTGGAAGAGATGAAGAGGGCAAAAGAGGATG 45 TGTTGCAGAAGGAGGTGAGAGTGAAGATTCTCAAGGACAACATCAAGTTATTAG CTGCCAAGGTGCCCTCTGGTGGCCAGGAGTTGACGTCTGAGCTGAATGTTGTGCT GGAGAATTACCAACTTCTTTGTAATAGAATTCGAGGAAAGTGCCACACGCTAGA GGAGGTCTGGTTGTTGGATTGAACTGCTTCACTATTTGGATCTTGAAACTACCT 

CGGATGCTGTCAACGAAGCCCTGGAGTCTCTGGAATCTGTTCTGCGCCACCCGGC CCTGGATGATAATCAGTGAGAAACTGGAGGCTTTCAACAGCCGATATGAAGA TCTAAGTCACCTGGCAGAGAGCAAGCAGATTTCTTTGGAAAAGCAACTCCAGGT 5 GCTGCGGGAAACTGACCAGATGCTTCAAGTCTTGCAAGAGAGCTTGGGGGAGCT GGACAAACAGCTCACCACATACCTGACTGACAGGATAGATGCTTTCCAAGTTCCA CAGGAAGCTCAGAAAATCCAAGCAGAGATCTCAGCCCATGAGCTAACCCTAGAG GAGTTGAGAAGAAATATGCGTTCTCAGCCCCTGACCTCCCCAGAGAGTAGGACT GCCAGAGGAGGAAGTCAGATGGATGTGCTACAGAGGAAACTCCGAGAGGTGTCC 10 ACAAAGTTCCAGCTTTTCCAGAAGCCAGCTAACTTCGAGCAGCGCATGCTGGACT GCAAGCGTGTGCTGGATGGCGTGAAAGCAGAACTTCACGTTCTGGATGTGAAGG ACGTAGACCCTGACGTCATACAGACGCACCTGGACAAGTGTATGAAACTGTATA AAACTTTGAGTGAAGTCAAACTTGAAGTGGAAACTGTGATTAAAACAGGAAGAC 15 CTTCCCTGAAGGTTCTTTACAATGACCTGGGCGCACAGGTGACAGAAGGAAAAC AGGATCTGGAAAGAGCATCACAGTTGGCCCGGAAAATGAAGAAGAGGCTGCTT CTCTCTCTGAATGGCTTTCTGCTACTGAAACTGAATTGGTACAGAAGTCCACTTC AGAAGGTCTGCTTGGTGACTTGGATACAGAAATTTCCTGGGCTAAAAATGTTCTG AAGGATCTGGAAAAGAGAAAAGCTGATTTAAATACCATCACAGAGAGTAGTGCT 20 GCCCTGCAAAACTTGATTGAGGGCAGTGAGCCTATTTTAGAAGAGAGGCTCTGC GTCCTTAACGCTGGGTGGAGCCGAGTTCGTACCTGGACTGAAGATTGGTGCAATA CCTTGATGAACCATCAGAACCAGCTAGAAATATTTGATGGGAACGTGGCTCACAT AAGTACCTGGCTTTATCAAGCTGAAGCTCTATTGGATGAAATTGAAAAGAAACC AACAAGTAAACAGGAAGAAATTGTGAAGCGTTTAGTATCTGAGCTGGATGATGC 25 CAACCTCCAGGTTGAAAATGTCCGCGATCAAGCCCTTATTTTGATGAATGCCCGT GGAAGCTCAAGCAGGAGCTTGTAGAACCAAAGTTAGCTGAGCTGAATAGGAAC TTTGAAAAGGTGTCTCAACATATCAAAAGTGCCAAATTGCTAATTGCTCAGGAAC CATTATACCAATGTTTGGTCACCACTGAAACATTTGAAACTGGTGTGCCTTTCTCT GACTTGGAAAAATTAGAAAATGACATAGAAAATATGTTAAAATTTGTGGAAAAA 30 CACTTGGAATCCAGTGATGAAGATGAAAAGATGGATGAGGAGAGTGCCCAGATT GAGGAAGTTCTACAAAGAGGAGAAGAAATGTTACATCAACCTATGGAAGATAAT AAAAAAGAAAAGATCCGTTTGCAATTATTACTTTTGCATACTAGATACAACAAA TTAAGGCAATCCCTATTCAACAGAGGAAAATGGGTCAACTTGCTTCTGGAATTAG ATCATCACTTCTTCCTACAGATTATCTGGTTGAAATTAACAAAATTTTACTTTGCA 35 TGGATGATGTTGAATTATCGCTTAATGTTCCAGAGCTCAACACTGCTATTTACGA AGACTTCTCTTTTCAGGAAGACTCTCTGAAGAATATCAAAGACCAACTGGACAAA CTTGGAGAGCAGATTGCAGTCATTCATGAAAAACAGCCAGATGTCATCCTTGAA GCCTCTGGACCTGAAGCCATTCAGATCAGAGATACACTTACTCAGCTGAATGCAA AATGGGACAGAATTAATAGAATGTACAGTGATCGGAAAGGTTGTTTTGACAGGG 40 CAATGGAAGAATGGAGACAGTTCCATTGTGACCTTAATGACCTCACACAGTGGA TAACAGAGGCTGAAGAATTACTGGTTGATACCTGTGCTCCAGGTGGCAGCCTGG ACTTAGAGAAAGCCAGGATACATCAGCAGGAACTTGAGGTGGGCATCAGCAGCC ACCAGCCAGTTTTGCAGCACTAAACCGAACTGGGGATGGGATTGTGCAGAAAC TCTCCCAGGCAGATGGAAGCTTCTTGAAAGAAAAACTGGCAGGTTTAAACCAAC 45 GCTGGGATGCAATTGTTGCAGAAGTGAAGGATAGGCAGCCAAGGCTAAAAGGAG AAAGTAAGCAGGTGATGAAGTACAGGCATCAGCTAGATGAGATTATCTGTTGGT TAACAAAGGCTGAGCATGCTATGCAAAAGAGATCAACCACCGAATTGGGAGAAA ACCTGCAAGAATTAAGAGACTTAACTCAAGAAATGGAAGTACATGCTGAAAAAC TCAAATGGCTGAATAGAACTGAATTGGAGATGCTTTCAGATAAAAGTCTGAGTTT

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ACCTGAAAGGGATAAAATTTCAGAAAGCTTAAGGACTGTAAATATGACATGGAA TAAGATTTGCAGAGGGGGCCTACCACCCTGAAGGAATGCATCCAGGAGCCCAG TTCTGTTTCACAGACAAGGATTGCTGCTCATCCTAATGTCCAAAAGGTGGTGCTA GTATCATCTGCGTCAGATATTCCTGTTCAGTCTCATCGTACTTCGGAAATTTCAAT GACCAGATGCTGAAGTCCAACATTGTCACTGTTGGGGATGTAGAAGAGATCAAT AAGACCGTTTCCCGAATGAAAATTACAAAGGCTGACTTAGAACAGCGCCATCCT CAGCTGGATTATGTTTTTACATTGGCACAGAATTTGAAAAATAAAGCTTCCAGTT CAGATATGAGAACAGCAATTACAGAAAAATTGGAAAGGGTCAAGAACCAGTGG GATGCCACCCAGCATGCGTTGAGCTAAGACAGCAGCAGCTTGAGGACATGATT ATTGACAGTCTTCAGTGGGATGACCATAGGGAGGAGACTGAAGAACTGATGAGA AAATATGAGGCTCGACTCTATATTCTTCAGCAAGCCCGACGGGATCCACTCACCA AACAAATTTCTGATAACCAAATACTGCTTCAAGAACTGGGTCCTGGAGATGGTAT CGTCATGGCGTTCGATAACGTCCTGCAGAAACTCCTGGAGGAATATGGGAGTGA TGACACAAGGAATGTGAAAGAAACCACAGAGTACTTAAAAACATCATGGATCAA TCTCAAACAAGTATTGCTGACAGACAGACGCCTTGGAGGCTGAGTGGAGGAC GGTGCAGGCCTCTCGCAGAGATCTGGAAAACTTCCTGAAGTGGATCCAAGAAGC AGAGACCACAGTGAATGTGCTTGTGGATGCCTCTCATCGGGAGAATGCTCTTCAG GATAGTATCTTGGCCAGGGAACTCAAACAGCAGATGCAGGACATCCAGGCAGAA ATTGATGCCCACAATGACATATTTAAAAGCATTGACGGAAACAGGCAGAAGATG GTAAAAGCTTTGGGAAATTCTGAAGAGGCTACTATGCTTCAACATCGACTGGATG TRESPONDE ATATGAACCAAAGATGGAATGACTTAAAAAGCAAAATCTGCTAGCATCAGGGCCC ATTTGGAGGCCAGCGCTGAGAAGTGGAACAGGTTGCTGATGTCCTTAGAAGAAC to grant TGATEAAATGGCTGAATATGAAGATGAAGAGCTTAAGAAACAAATGCCTATTG GAGGAGATGTTCCAGCCTTACAGCTCCAGTATGACCATTGTAAGGCCCTGAGACG GGAGTTAAAGGAGAAAGAATATTCTGTCCTGAATGCTGTCGACCAGGCCCGAGT TTTCTTGGCTGATCAGCCAATTGAGGCCCCTGAAGAGCCAAGAAGAAACCTACA ATCAAAAACAGAATTAACTCCTGAGGAGAGAGCCCAAAAGATTGCCAAAGCCAT GCGCAAACAGTCTTCTGAAGTCAAAGAAAAATGGGAAAGTCTAAATGCTGTAAC TAGCAATTGGCAAAAGCAAGTGGACAAGGCATTGGAGAAACTCAGAGACCTGCA GGGAGCTATGGATGACCTGGACGTGACATGAAGGAGGCAGAGTCCGTGCGGAA TGGCTGGAAGCCCGTGGGAGACTTACTCATTGACTCGCTGCAGGATCACATTGAA AAAATCATGGCATTTAGAGAAGAAATGCACCAATCAACTTTAAAGTTAAAACGG TGAATGATTTATCCAGTCAGCTGTCTCCACTTGACCTGCATCCCTCTCTAAAGATG TCTCGCCAGCTAGATGACCTTAATATGCGATGGAAACTTTTACAGGTTTCTGTGG ATGATCGCCTTAAACAGCTTCAGGAAGCCCACAGAGATTTTGGACCATCCTCTCA GCATTTTCTCTCTACGTCAGTCCAGCTGCCGTGGCAAAGATCCATTTCACATAAT AAAGTGCCCTATTACATCAACCATCAAACACAGACCACCTGTTGGGACCATCCTA AAATGACCGAACTCTTTCAATCCCTTGCTGACCTGAATAATGTACGTTTTTCTGCC TACCGTACAGCAATCAAAATCCGAAGACTACAAAAAGCACTATGTTTGGATCTCT TAGAGTTGAGTACAACAAATGAAATTTTCAAACAGCACAAGTTGAACCAAAATG ACCAGCTCCTCAGTGTTCCAGATGTCATCAACTGTCTGACAACAACTTATGATGG ACTTGAGCAAATGCATAAGGACCTGGTCAACGTTCCACTCTGTGTTGATATGTGT CTCAATTGGTTGCTCAATGTCTATGACACGGGTCGAACTGGAAAAATTAGAGTGC AGAGTCTGAAGATTGGATTAATGTCTCTCCCAAAGGTCTCTTGGAAGAAAAATA CAGATATCTCTTTAAGGAAGTTGCAGGGCCAACAGAAATGTGTGACCAGAGGCA GCTGGGCCTGTTACTTCATGATGCCATCCAGATCCCCCGGCAGCTAGGTGAAGTA GCAGCTTTTGGAGGCAGTAATATTGAGCCTAGTGTTCGCAGCTGCTTCCAACAGA ATAACAATAAACCAGAAATAAGTGTGAAAGAGTTTATAGATTGGATGCATTTGG

AACCACAGTCCATGGTTTGGCTCCCAGTTTTACATCGAGTGGCAGCAGCGGAGAC TGCAAAACATCAGGCCAAATGCAACATCTGTAAAGAATGTCCAATTGTCGGGTTC AGGTATAGAAGCCTTAAGCATTTTAACTATGATGTCTGCCAGAGTTGTTTCTTTTC GGGTCGAACAGCAAAAGGTCACAAATTACATTACCCAATGGTGGAATATTGTAT 5 ACCTACAACATCTGGGGAAGATGTACGAGACTTCACAAAGGTACTTAAGAACAA CAGACAGTTCTTGAAGGTGACAACTTAGAGACTCCTATCACACTCATCAGTATGT GGCCAGAGCACTATGACCCCTCACAATCTCCTCAACTGTTTCATGATGACACCCA TTCAAGAATAGAACAATATGCCACACGACTGGCCCAGATGGAAAGGACTAATGG 10 GTCTTTCTCACTGATAGCAGCTCCACCACAGGAAGTGTGGAAGACGAGCACGCC CAGAGCCCAGCTCAGATCCTGAAGTCAGTAGAGAGGGAAGAACGTGGAGAACTG GAGAGGATCATTGCTGACCTGGAGGAAGAACAAAGAAATCTACAGGTGGAGTAT GAGCAGCTGAAGGACCAGCACCTCCGAAGGGGGCTCCCTGTCGGTTCACCGCCA 15 GAGTCGATTATATCTCCCCATCACACGTCTGAGGATTCAGAACTTATAGCAGAAG CAAAACTCCTCAGGCAGCACAAAGGTCGGCTGGAGGCTAGGATGCAGATTTTAG AAGATCACAATAAACAGCTGGAGTCTCAGCTCCACCGCCTCCGACAGCTGCTGG AGCAGCCTGAATCTGATTCCCGAATCAATGGTGTTTCCCCATGGGCTTCTCCTCA GCATTCTGCACTGAGCTACTCGCTTGATCCAGATGCCTCCGGCCCACAGTTCCAC 20 CAGGCAGCGGAGAGCCTGCTGGCCCCACCGCACGACACCAGCACGGATCTC ACGGAGGTCATGGAGCAGATTCACAGCACGTTTCCATCTTGCTGCCCAAATGTTC GTACAGTGTTGCCCTTTTCAGCAAATGCCAATTCCAAGTTCCATTAAATCAGAAG... CECCATGGCTCCTTGGCCCACGATGTTGAGTGCTGACTGTGTTCTACTGAAAG 25 Á ÁGTÁAAACACTGACTATCCAAAGAGAAATGGATATTTTGTTTTTATAATAACCAT ATATTATTGTTTCTTCTCCCTTTCTATGCAAGTGTAAATTAATGAACAGAGAGG TATTTGGAAATGGTAATACATTTGTCACGGATTTGTATAATGTATACAGCATTGG GAAAGTGGGTGGGGCTTTCTAATATGATACCGTCTTTTTAATAACTATGACAAA GCTTACATAAGAATTAGAAGACCACTTTACATTTTTACATTCCTTCTGCTGTTCAT 30 ATTAACCTTGCACAATTACTTCATTTTTCTTTGACTCTTTTACCACAATGTTTTGG TTATTATAATTTATCAGCCATATGTTTATCAGCCATATAACCAACTAGATCCCAA ATAGATCCATGTATTTGTTTCCGTGATTTGGCCACATTAATAAATTCATAAATTTC AATCAAATATCATATATACACACATATGGTTTAAGCTACAGCCCTGTGTATGC CGTTTAACTTTATTTGACGTTGCCCACTTACTTCTTTGCTGACCACTTGGATAACC 35 GTAATAAAAATCCTATAAGCCTAAATGGCATTTCTTTTGGGATATTTTTCCTGCAT TGATAAAGAAGACTACATTATAATAATCTCAAAGATCATATTACCAAAGGTTGCC CACTTGAGCATATTTCATTTTGACACAGAAACAAAATTTAGTACAACCTTTCCT AGTTCCCATGTCTTGATTTCATCATTACATGCACAGCAGACCTTTACCTATTGTG 40 ATACCAGAACACATCATTGTCTTTGGTTCCCTTCAAAGAGAATTTTATTGTTGTTT TGTATTTCAAGTCCTTAATAGTTCTTGAAACTCCTAGTTGTTTCTTGTTGAAAG CAGACACACATTTAGTGCACGGCTTATTTTACCTTTCGGGTGAAAGATCAGATGT TTTTATACCCTTCACTTGATCAATATATTTGGAAAGAATGTTTATCAAAAGTCTAT GTCACTGCTTCTACAGAAGAATGAAATTAATGCTTAGGTGATGGTACCTCCACCT 45 ACATCTTTTGAGTGCATTCAATTATGTATTTTGGTTTAGCTTCTGATTTAACATTT AATTGATTCAGTTTAAACATGTTACTTAATTAGCAAATGTAGAGGAACCAAAAAA AGGTGAAAATAATATGTTTTGATTCAAACCTAAAGACATAAAAACATAAAGACA TTTTAACTTTGGGTTCTCTTTAGCTGGGATCTGGCCAGAAGGAGGCTTAAAGTTA GAAATTGCTATTATTTTAGAATAGGTTGGGTGGGTTGGGGGGCAAGGGTGTCTAT

TTGCAGCAGAGATATTTTGAAAAGAAGAAAATTGTTTTATATAAAAAGGAAAGC CATGACCACCTTTCTACCTCAGATCCATCTTCATCCATTGCATTGGAAACTGCTTT ATGCTGCTGCAGTCTGCAAAGTCTAGAGCTTTTATCAGGCCATGTCATACCCAAG AAAGCACCTATTTAAAGAAAAAACAATTCCCTGAGCTCTCAACTCCAAGTTGTAG 5 TGTATTGTATGCAAAATGTCCTCTATCTGCTATTAAAGAAAAGCTACGTAAAAC ACTACATTGTAACCTTCTAAGTAATAATAAATAAAAAGAAATATATTGCAGTAAC AATGGGAAGTAAGTATGTAGTTCTTTTGAAATATGTGGTAAAGAACTAATCACAG ACTATCATCTAATCTGGTTACATATTGTATTTTCATCCTGTGATTAAAAGGCACA 10 TGTGTAAAAGTCCAATTAGTATGCTTTTCATTTCAAATAATCCATATAGCCTCCAG GTATTCTGTATTTGTATAAAGTACGTGCAAACACCTTTCTGCTAATCGGGTCCCC ACATTCTTTCACTACAGGTACTTTACAAGTCTGCCCTCTGCTCAAACACTAACCG 15 GACATCCTCCTGACCTCCCCTGACCTGCTTCACCACTGTGTTACCTCA CTGGTTACTTGTTACAGCAAACTGATGCAACTACTAGTCTACCTGGACAACATAT TAAACAGGTATCACCTAATAGGGTGGCAGCCTATCGGGGTGATTCCTGGCGAAT ACACAGTAACCAACCACATACTGACACACTCAACCCATTTGCTACAGATGGACCC ACACTAATTGATATGACAATCCTTTATTCACTCGGCACATTTGGTTTCTTTGCATT 20 TTCTTCCATTTTACATTGCAGGTGTGGCTACCAAGAGCTGGATAACGAGTCCCTC AAACAAAGTTTGGAATTGCGAGATATATTGGGGTACCTTGATTCTTGAGACAGTT TO THE GRANT CITETAGT TGATCTTGT TGTATTAAAAAGTCACTCTCAACTGAAGTGACC ACTGCATTTCTTTTGTAAAAAAGGTCATTTGACTGGCTTTTCCTCACAACTGCCACC **T25** ACATTTCACATTTTTTAAAAAAAAAAACAATCCTTCATGGGAATATATCCTAATAATC AATTATATGGAGACAGTTTTATGTACACCAAATTTCTGCAACTTTATAATAATGA TTGCATCCATTAAATGGAATATAATATAGCCATTAAAATTATGTTTTTGTAAAATT TTTAATGCCATAAGAAAATGTGGCAATTTTGCAATGAAAAAGATCTACTTATAAA 30 ACTGTTTACAGTATGACTCCAATTATGTAAAAAAAGTATACAATACACATATAGG CATACATGGGGGTTGCTTTTTAAAGGTGGTTACTTCTGGGTTGTGATATTATCAGT AATCATTTTTGCTTTTTTATACATTTCTGTATTTTCAAGTTTTCTATGATGAGTAT ATTATTTTACAAAGACTACGAAAATTTTCCTCTGATATACTGGTAATTAGAATGT ACTTGGGTATTTTAAATATGGGAACAATATTATAGTGCTTCATCTTCTATGACT 35 TTTTTGGAATACATATCACTTTGGTAATAAACTTACATTCCCTGTTTTATACTTGT TACAACATTTAATTAAACAGTTAATATTGTGATTAGAGCATTGTTTGCTTCATGA CCTAAACAAATACTGGCTTTGAAGTCTAGGTTCTATTTCCTAGAAGATTTAACAT GCTTTCTAAACAAAAGATAATTCCAACTTACAGTTTTCCTATGTAAGGGAAAAAA 40 ATAGGCATAAACGTGTTTATTAAGTGAAACGTATCCTTTAAAAAATAAAAAAGGG AAGCCTGTATATAAATGAAGTTGTGGATTCAACTAGCCAGAATTTATTCTGACTT GCACCAAACCACAAAATCTTTTAAAAGTCTAGTTAGTGTAGTCTAAATGGACA CTCCAGAGTCTGTTCTTGAATTCCATTGCAAGAGCTCCAACTTCCTACTTTCAGAA 45 GGGATGGGGATCAAGATGAGGGTTGTCACATAAGCTAATTTTCAATATATCAA GTCTTGTGGGGTCCAGGAACAAATACTGTCATTGGTTAGTGTTTAAGTACATGAG TTGACTTTTCTCCTCTCACACCCCACCTTGCCCTGGCAATTGGGTAGGGGGAG GCTGTTTATCCTCCAAGAGAGGCGGCTGGTTCCTCATCTCAGTTTCCGTTCTAAA CCACAGAGTGGTCATTGCTGTGAACTCCAGCCAAGATGGTGTGGGAGAGGCGAG

GAAGCCGAGCGTCTGAGCCTTCTGTGGGGCCGGTGGGGTTCTCACTGCGCTGGC AGCAGAGGATCTGCCTAAAGGTGGCGCTCATTTCTTTGTCGCGGTAGGAGTAAAT GGCCAGCACGTCGCACTGTGGACAGCACACGTCTAGAAGTAACAAAACCAATCC 5 AGGAGTCCAGCAGATGATAAAGGCCCCAAGCACAATGACCACAGTCTTCAGAAG ACTCATCATGGTATCCCGATTCCGCCGGGGTCCAGAACTATGCCGAGACATTCTC ATAGTCCTCTGGCGAACATAGCCAAAGATGTGAGCATAGAGAACCACCATTACC AGGGGTGCCATGTTGGAACAATTTTCAATATCACAGATACAGTTCCAGCCCACAC 10 TGGGTATAGCACCCATAACGATGGCCATAGTCCAGATGACCACAATGACCACCA CTACCGCCGGTTGCTCATCCGTGTGTGGAGCTGCATGCGGAAAACCGTAATGTG CCTCTCGATTGCAATAGCCAGTAAGTTGGCCACAGATGCCGTCAGGCTGGTGTCA CCTGTGTTGAACATGAGATAGAAGTAGGCCAACCCAGCAAAGAAGTCTGCAGCA 15 GCCAGATTAGCCATTAGGTAATAAATAGGAAAATGGAAGCGGCGGTTGACATAG ATTGCCACCATGACCAATAGGTTGGCCAACATGATGAAGATACAAACAGTGATT CCAAGTCCCATCACCAGCTTGCTGACTGTGTTCCATTCTGTGGCAAGATGCTTTCC ACTTCGGTTATAAAAGAAGGCAATGGACTCGTTGTAGAAGCACTGTGGTTCATTC ATGGCTGTGAACTGGGGCTGTGAAATTACAGGGATGGAAGTAGAGATGGCAGCC 20 ATGACAGCTCTGTGGTTGTAGGTGGTGAACACGCCCCAGAACTACGGGAGACAA ATTTCTTGTTTGCTGATCAGATCGAAGTCATGCTAGGAGAAGCTGTGTACCTGA AND AND STATEMENTAL AND ACCIDENCE OF THE STATEMENT OF THE VIII I CAGAAATCCATGCTGAGTGCCCACAGACCTGGGCAGGAGCTGTTCCCCCAGCGCCG 25 GACAGCTGGCAGGACTCCGGTGGACGCCCCGGCACGGGCATTTTCACGTTGTC GCTCTCCTCTTCCCACTTGAAAAGCTCTGGAAAACATCGCGGGGCCCGCAAAACC CCGGAAATGTGGC

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yc03e09.s1 Stratagene lung (#937210) Homo sapiens cDNA clone IMAGE:79624 3', mRNA sequence gi|666284|gb|T62627.1|T62627[666284]

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AATTCACTTGCTATNTAACTCTCTCTNGAGATTTATTCTTGGAGGACAGAGCAAA
AGTCCACTCTTCAGCAGCTCTCCGAGGGTCATTCCTTCACAACGTATATTCCGTTT
CCAGTTCTTTGCGTTCCTTCCTTTTCCTTCGACTTCAAATTCATTTGGTGTTAACCA
AGTTCCATCCTCATTCCNGAATGCACTTCACTGAGGATCCCGTGTTTCATTTTCTT
CTTATATAAAANCCCTTTCGCCTCACCACAGGTCACGGGGGAGCTTNGGAACAGT
GAAAATCCACAGTGTCACTTTGGGGTTTTCCTCTTTGAA
ATCTCCTTTTTGAGCTTGGACAGATATCTTGNTCCTTTTGNCT

40

**SEQ ID NO: 671** 

ys88a08.s1 Soares retina N2b5HR Homo sapiens cDNA clone IMAGE:221846 3' similar to SP:HTLF\_HUMAN P32314 HUMAN T-CELL LEUKEMIA VIRUS ENHANCER FACTOR; contains MER22 repetitive element;, mRNA sequence

45 gi|1064703|gb|H84982.1|H84982[1064703]
GCTCCCAGTGGTCAGCGGAGACCCCAAGGAGGATCACAACTACAGCAGTGCCA
AGTCCTCCAACGCCCGGAGCACCTCGCCCACCAGCGACTCCATCTCCTCCTC
CTCCTCAGCCGACGACCACTATGAGTTTGCCACCAAGGGGAGCCAGGAGGCAG
CGAGGGCAGCGAGGGGAGCTTCCGGAGCCACGAGAGCCCCAGCGACACGGAAG

AGGACGACAGGAAGNACAGCCAGAAGGAGCCCAAGGATTTTTTNGGGGACAGC **GGGTACGATTNCC** 

**SEQ ID NO: 672** 

- yq55b04.r1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:199663 5' 5 similar to SP:SISD HUMAN P13501 T-CELL SPECIFIC RANTES PROTEIN PRECURSOR;, mRNA sequence gi|982328|gb|R96668.1|R96668[982328] NCGCCCAGGAGTCCTCGGCCAGCCCTGCCCACCAGGAGGATGAAGGTCTC CGTGGCTGCCTCTCCTGCCTCATGCTTGTTGCTGTCCTTGGATCCCAGGCCCAGT
- 10 TCACAAATGATGCAGAGACAGAGTTAATGATGTCAAAGCTTCCACTGGAAAATC CAGTAGTTCTGAACAGCTTTCACTTTGCTGCTGACTGCTGCACCTCCTACATCTCA CAAAGCATCCCGTGTTCACTCATGAAAAGTTATTTTGAAACGAGCAGCGAGTGCT CCAAGCCAGGGTGTCATATTCCTCACCAAGAAGGGGCGGCAAGTCTGTGCCAAA CCCAGTGGGTCCGGGAGTTCAGGATTGGCATGGAAAAAGCTTNAAGCCCTAATT
- 15 CAATATTANTAATTAAAGGAGGACANAAGAGGGCCAGCNCACCCACCTCCAACA CTTCNTGAGGCTTTGGAAGG

**SEQ ID NO: 673** 

zt20b07.s1 Soares ovary tumor NbHOT Homo sapiens cDNA clone IMAGE:713653 3' similar to TR:G577291 G577291 MRNA; contains element MER28 repetitive element; 20 mRNA sequence

- CCGCCTCCTTTGCCGGGGTACACCTGGCCCACAAGAGACCTTCAGCACGTGTCGA \*\*\*\*\*\*\*CTTCTCAAAGATAGACEEGGGCATAGCCTGAAAGCATATTGAAAATGACGAAAA 25 AAGGGAAGACTCTCATGATGTTTGTCACTGTATCAGGAAGCCCTACTGAGAAGG AGACAGAGGAAATTACGAGCCTCTGGGAGGGCAGCCTTTTCAATGCCAACTATG
  - ACGTCCAGAGGTTCATTGTGGGATCAGACCGTGCTATCTTCATGCTTCGCGATGG GAGCTACGCCTGGGAGATCAAGGACTTTTTGGTCGGTCAAGACAGGTGTGCTGAT GTAACTCTGGAGGGCCAGGTGTACCCCGGCCAA GGAGGAGGAA

30

**SEQ ID NO: 674** 

>L01639

CGCATCTGGAGAACCAGCGGTTACCATGGAGGGGATCAGTATATACACTTCAGA TAACTACACCGAGGAAATGGGCTCAGGGGACTATGACTCCATGAAGGAACCCTG

- 35 TTTCCGTGAAGAAATGCTAATTCAATAAAATCTTCCTGCCCACCATCTACTCC ATCATCTTCTTAACTGGCATTGTGGGCAATGGATTGGTCATCCTGGTCATGGGTT ACCAGAAGAAACTGAGAAGCATGACGGACAAGTACAGGCTGCACCTGTCAGTGG CCGACCTCCTCTTGTCACACGCTTCCCTTCTGGGCAGTTGATGCCGTGGCAAACT GGTACTTTGGGAACTTCCTATGCAAGGCAGTCCATGTCATCTACACAGTCAACCT
- 40 CTACAGCAGTGTCCTCATCCTGGCCTTCATCAGTCTGGACCGCTACCTGGCCATC GTCCACGCCACCAACAGTCAGAGGCCAAGGAAGCTGTTGGCTGAAAAGGTGGTC TATGTTGGCGTCTGGATCCCTGCCCTCCTGCTGACTATTCCCGACTTCATCTTTGC CAACGTCAGTGAGGCAGATGACAGATATATCTGTGACCGCTTCTACCCCAATGAC
- 45 TATTGTCATCCTGTCTGTATTGCATTATCATCTCCAAGCTGTCACACTCCAAGG GCCACCAGAAGCGCAAGGCCCTCAAGACCACAGTCATCCTCATCCTGGCTTTCTT CGCCTGTTGGCTGCCTTACTACATTGGGATCAGCATCGACTCCTTCATCCTCCTGG AAATCATCAAGCAAGGGTGTGAGTTTGAGAACACTGTGCACAAGTGGATTTCCA TCACCGAGGCCCTAGCTTTCTTCCACTGTTGTCTGAACCCCATCCTCTATGCTTTC

**SEQ ID NO: 675** 

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> Human tumor necrosis factor receptor 2 (TNFR2) gene, exon 10 and complete cds 10 gi|1469539|gb|U52165.1|HSTNFR2S10[1469539] TCTTGGTCTCGGCTCCTGGCCCAGTGCTCTTTCCCATGTGTCTGAATCTGCATCTT GGGCAGGGGTCCCTGGGCCCCACTCCTGGACCCCGGACTGACCCCCACCCCATC TTGTGCTTAGCAGATTCTTCCCCTGGTGGCCATGGGACCCAGGTCAATGTCACCT GCATCGTGAACGTCTGTAGCAGCTCTGACCACAGCTCACAGTGCTCCTCCCAAGC 15 CAGCTCCACAATGGGAGACACAGATTCCAGCCCCTCGGAGTCCCCGAAGGACGA GCAGGTCCCCTTCTCCAAGGAGGAATGTGCCTTTCGGTCACAGCTGGAGACGCCA GAGACCCTGCTGGGGAGCACCGAAGAGAAGCCCCTGCCCCTTGGAGTGCCTGAT GCTGGGATGAAGCCAGTTAACCAGGCCGGTGTGGGCTGTGTCGTAGCCAAGGT GGGCTGAGCCCTGCAGGATGACCCTGCGAAGGGGCCCTGGTCCTTCCAGGCCC 20 CCACCACTAGGACTCTGAGGCTCTTTCTGGGCCAAGTTCCTCTAGTGCCCTCCAC AGCCGCAGCCTCCCTCTGACCTGCAGGCCAAGAGCAGAGGCAGCGAGTTGGGGA A TOWN COLOR COLOR TO THE TERM OF THE TERM 25 TTTGTTTCTCCCCTGGGCTCTGCCCAGCTCTGGCTTCCAGAAAACCCCAGCATCC TTTTCTGCAGAGGGCTTTCTGGAGAGGGAGGGATGCTGCCTGAGTCACCCATGAA GACAGGACAGTGCTTCAGCCTGAGGCTGAGACTGCGGGATGGTCCTGGGGCTCT GTGTAGGGAGGAGGTGGCAGCCCTGTAGGGAACGGGGTCCTTCAAGTTAGCTCA GGAGGCTTGGAAAGCATCACCTCAGGCCAGGTGCAGTGGCTCACGCCTATGATC 30 CCAGCACTTTGGGAGGCTGAGGCTGGATCACCTGAGGTTAGGAGTTCGAGA CCAGCCTGGCCAACATGGTAAAACCCCATCTCTACTAAAAATACAGAAATTAGC CGGGCGTGGTGGCGGCACCTATAGTCCCAGCTACTCAGAAGCCTGAGGCTGGG AAATCGTTTGAACCCGGGAAGCGGAGGTTGCAGGGAGCCGAGATCACGCCACTG 35 GCACCGCCTCCAAATGCTAACTTGTCCTTTTGTACCATGGTGTGAAAGTCAGATG CCCAGAGGCCCAGGCAGCCACCATATTCAGTGCTGTGGCCTGGGCAAGATAA CAACAAGCCAACGACAAACTCTGCCAGCCACATCCAACCCCCACCTG 40 TGCTGTCCTAGGCCACACCATCTCCTTTCAGGGAATTTCAGGAACTAGAGATGAC TGAGTCCTCGTAGCCATCTCTCTACTCCTACCTCAGCCTAGACCCTCCTCCTCCCC CAGAGGGGTGGGTTCCTCTTCCCCACTCCCACCTTCAATTCCTGGGCCCCAAAC GGGCTGCCCTGCCACTTTGGTACATGGCCAGTGTGATCCCAAGTGCCAGTCTTGT GTCTGCGTCTGTGCGTGTCGTGGGTGTGTAGCCAAGGTCGGTAAGTTGAA 45 TGGCCTGCCTTGAAGCCACTGAAGCTGGGATTCCTCCCCATTAGAGTCAGCCTTC CCCCTCCAGGGCCAGGGCCCTGCAGAGGGGAAACCAGTGTAGCCTTGCCCGGA TTCTGGGAGGAAGCAGGTTGAGGGGCTCCTGGAAAGGCTCAGTCTCAGGAGCAT 

AATTGTTGATAAATTCCACTGGACTTGAGCTTGGCAGCTGAACTATTGGAGGGTG

10

15

SEQ ID NO: SEQ ID NO: 676

>R88734

TGACTTTTAAAAACTGACATTGTATTGAATTTACATAATTCTCAATCAGAAAAAA AATTACTGTCAGACTGCAATGCA AGTCTGCCCCAATGAAGGCCG

20 SEQ ID NO: 677 >AA418689

- 25 TCTTTCGGGCCTTGAGTTCCTTCATGGCAATGAGCAGAGGATCTGTCTCCCCCT©C
  AGCTCCACCATCACAGGGGCACACATCGCAATCTGGAGCGCTCGGGTGCCCAGC
  ACGCGGGCTCGCTCGTACTTGGTCATGTATGGTGGTGATTCGCTTCTGGTTGG
  CCTGCGGTCGCTCCCCAGAGGAGGNAGTCTCGACATTCTCCTGGCCTTCCTCTC
  GGCATTCTCCAAGTCATCTAGCCCTTCATCCTCCTC
- 30 CACATCATCAGAGTCGTCGCCATCAAA

SEQ ID NO: 678 >AA455281

TTTTTGGAGGAGTGGCATGGAGTTCTTTAATTTGGAAGGCAAAAGGTTACATTTA

ATGAAAGGCAGAGGCTGGATTAATAAATGTTTGTTAGAAAAGTTGTTCTGACACAC
AGTGAACTCTGGGCTTTTCTCCTGCATAAAAAAGCAGAGCTAGCAGTAAGTGCAA
ATCTGAAGAAAATCCATGTGTCCAATAAGCTGCCATCTCCAGAACTCTTATCCAG
GAAATTCAAAGAGTGAACATTCTTTTAGTCTCCTACTCCTCAATTAAGTAAATGA
GAATGAGTCAGCCAACAAAGTTCATGACAACAAGGTGCAGGATGGTGCTGGCAA

40 AGAGAAATCAGCAAAGGCTCGCTCTGGGGAGATGCCTTGGAAATCCGCTTTGT TCTGTGGGTTGATCTGTATTCTCAGGCAAACCGCTAGGATGAAACTCCCCACACA AGAGATGAAGCCCGAGAGAAAAGAGTTGAAGGGGAAGGTCCC

**SEQ ID NO: 679** 

45 >H94469

GCAAAACAACATTTATTCTTTTAAAAAAATCTATATACATTGCCATACAAAGATAC CACATTGAAGCAGTTCTCAGGAACCTTCCAGTGAGCCTTCTTATAATTGCCCG AGCAAGATTTCGTGCCAGAGAAAGTCTCAGCATTTCCACCTTGGTGTNCTCTATG TCATCATCCTGGAGCTGCTCGGTATCAGATTCTCCATGCACAGGTCTTCTTGACGT

CAAGTCCTCCAGACACCGCATCAACTCATAAGTCTGTTCTGCTGAGAAAATCACC TGTTTCTGTTCCAAAAGGGGCAAGGCATCTGTCAGCAGAGTTCATCCCAGAAAGA CCGAAGGGGCAATCCGAGACGTCATCAAG GACAGAAGGA

- 5 SEQ ID NO: 680
  - aa79c05.s1 NCI\_CGAP\_GCB1 Homo sapiens cDNA clone IMAGE:827144 3' similar to SW:RLX1\_HUMAN P49406 PUTATIVE 60S RIBOSOMAL PROTEIN;, mRNA sequence gi|2261786|gb|AA521243.1|AA521243[2261786] TTTTTTTTTGGTGTACAAGTTTTATTTTAGAAAAAAAGTATTAATAAAACAATGA
- 15 AAATTTGGACGTTCCCAGCGTTTAGACCAGGGCTTAGGCTTCATTTTACTTTCAG CTCATTAACAGGAACTTTTTGGTTAGGCTCTTGTACTACTGGCTTCATATTCACAT CAAAAGTGCTATATTCAGGAAGGGCATCTCGTAAGTATAGCAAGCTATCATCCA GCCGTTTCTCTAATTTGACCACCTGAATCTCCTGGACCCGAGGATTATAAAGTTC AAAGCAAATCTCGACACCTTGTCCTTCGATAACATTCCTAAGGAT GAAAGTAGC

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**SEQ ID NO: 681** 

- in Lives Human Thy-1 glycoprotein gene, complete cds (1000 per section) and the section of the section of the complete cds (1000 per section) and the section of the sectio
  - GGATCCAGGACTGAGATCCCAGAACCATGAACCTGGCCATCAGCATCGCTCTCCT
  - 25 GCTATCAGGTACCCGGCATGGGGCAGGACTGGGGTCCCAGGCGCCCTGGCTTCCC TTCCCTCCAGAGAAGCAGCTTCTCCCTCACAGTCTCAGAAAAGCGCAGGTGACAA AGAGAGGGCTCTTTTCATCCTGAAGTCAGCCGATCCACCGCGCTGATATTCTGA CGGCCTGAGGTGGTTTTTGGAAACACAGTTTGCTGAGCCCTCCTTCACACTATTG AACTAGAATCCCCAACTGAGAACCCAGGAACCAGCATCAACTCCCTAAGATCTC
- 35 GTGGACCAGAGCCTTCGTCTGGACTGCCGCCATGAGAATACCAGCAGTTCACCCA TCCAGTACGAGTTCAGCCTGACCCGTGAGACAAAGAAGCACGTGCTCTTTGGCAC TGTGGGGGTGCCTGAGCACACATACCGCTCCCGAACCAACTTCACCAGCAAATA CCACATGAAGGTCCTCTACTTATCCGCCTTCACTAGCAAGGACGAGGGCACCTAC ACGTGTGCACTCCACCACTCTGGCCATTCCCCACCCATCTCCCCAGAACGTCA
- 45 AACAGGATGACACCACCTCCCTCAGCCAGTTTTCTTGTCATGATGTTTAGTAAG GTTTTCATAAGATGATATGTGTGCAAGAGATCAGTAATCTGCAAATGGGAAAGA TGGCTGGTTCTGTGAGACCAGGCTGTTCCTGGTCCCAGCTAAGACATTGCAGTAC CCACCTCCCAAAGGGAGTACACCCTTGCTTTGGGCCTGTGCCTGAGTCCTG ATCCGTCTTCCTTCCTACCCTGCCCCCGGCCCCCTTCTCTTTCTGCAGACAAACTG

GTCAAGTGTGAGGCATCAGCCTGCTGGCTCAGAACACCTCGTGGCTGCTGCTGC TCCTGCTGTCCCTCCCTCCAGGCCACGGATTTCATGTCCCTGTGACTGGTG GGGCCCATGGAGGAGACAGGAAGCCTCAAGTTCCAGTGCAGAGATCCTACTTCT CTGAGTCAGCTGACCCCCTCCCCCAATCCCTCAAACCTTGAGGAGAAGTGGGGA 5 CCCCACCCTCATCAGGAGTTCCAGTGCTGCATGCGATTATCTACCCACGTCCAC GCGGCCACCTCACCCTCTCGCACACCTCTGGCTGTCTTTTTGTACTTTTTGTTCC TGAAGAGGGAAGCCAGGATTGGGGACCTGATGGAGAGTGAGAGCATGTGAGGG GTAGTGGGATGGTGGGTACCAGCCACTGGAGGGGTCATCCTTGCCCATCGGGA 10 CCAGAAACCTGGGAGAGACTTGGATGAGGAGTGGTTGGGCTGTGCTGGGCCTAG GACCCCAGATGTGAGGGCACCACCAAGAATTTGTGGCCTACCTTGTGAGGGAGA GCCCTCCTTACCACTGTGGAAGTCCCTCAGAGGCCTTGGGGCATGACCCAGTGAA 15 GATGCAGGTTTGACCAGGAAAGCAGCGCTAGTGGAGGGGTTGGAGAAGGAGGTA AAGGATGAGGGTTCATCATCCCTCCCTGCCTAAGGAAGCTAAAAGCATGGCCCT GCTGCCCCTCCCTCCACCCACAGTGGAGAGGGCTACAAAGGAGGACAAGA CCCTCTCAGGCTGTCCCAAGCTCCCAAGAGCTTCCAGAGCTCTGACCCACAGCCT CCAAGTCAGGTGGGGTGGAGTCCCAGAGCTGCACAGGGTTTGGCCCAAGTTTCT 20 TGAGCCCTCAGACAGCCCCTGCCCGCAGGCCTGCCTTCTCAGGGACTTCTGC → #\###GGGGCCTGAGGCAAGCCATGGAGTGAGACCCAGGAGCCGGACACTTCTCAGGAA ######ATGGCTTTTCCCAACCCCAGCCCCACCCGTGGTTCTTCCTGTTCTGACTGT ATAAAACCAAGCCTCTGGAATCTGTCCTCGTGTCCACCTGGCCTTCGCTCCTCCA 25 GCAGTGCCTGCCTGCCCCGCTT

**SEQ ID NO: 682** 

yw08h11.s1 Soares melanocyte 2NbHM Homo sapiens cDNA clone IMAGE:251685 3',

30 mRNA sequence gi|1110224|gb|H96738.1|H96738[1110224]
TAAAANAAATCTTTTTTTATTTCAAAGATTGCTTCTTATATTGAAGCTCATATTA
AAGCAACAGTACAATGTTCATAAAATATAAGTGTGATGCCGTAACATTTTCTTAC
ATGTCAGAATACTGATATTTATATGTATACTAAAATAAGAACTTTAAAATTGTAC
AAATAGATACATTAAAAATGACATAGAAATAGGGCGTCTCTCACTGAAACAAGA

35 CAGTTATATCTGGCACGTATTAGTTTAAGATGAAAGTAGAAGCAAAAAGATTTAC AAGAATCAGCAGTAACAAGATTGATGCTCAAGAGACATAATTGTACATTGTATT GTACATACATTGTATGGGTTTAAGCTGGCTGGAATATTATATATTTCCAAGTTTTA AAAATGGCNCTACCANATAGAGTGGTCCNGAGTTTAAGGCGAAATTACAGCTCA GAACTGTTGTCCCTTCNAATTTTGGTGG

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**SEQ ID NO: 683** 

Human integral membrane serine protease Seprase mRNA, complete cds gi|1924981|gb|U76833.1|HSU76833[1924981]

CCACGCTCTGAAGACAGAATTAGCTAACTTTCAAAAAACATCTGGAAAAATGAAG
45 ACTTGGGTAAAAATCGTATTTGGAGTTGCCACCTCTGCTGTGCTTATTGGT
GATGTGCATTGTCTTACGCCCTTCAAGAGTTCATAACTCTGAAGAAAATACAATG
AGAGCACTCACACTGAAGGATATTTTAAATGGAACATTTTCTTATAAAACATTTT
TTCCAAACTGGATTTCAGGACAAGAATATCTTCATCAATCTGCAGATAACAATAT
AGTACTTTATAATATTGAAACAGGGCAATCATATACCATTTTGAGTAATAGAACC

ATGAAAAGTGTGAATGCTTCAAATTACGGCTTATCACCTGATCGGCAATTTGTAT ATCTAGAAAGTGATTATTCAAAGCTTTGGAGATACTCTTACACAGCAACATATTA CATCTATGACCTTAGCAATGGAGAATTTGTAAGAGGAAATGAGCTTCCTCGTCCA ATTCAGTATTTATGCTGGTCGCCTGTTGGGAGTAAATTAGCATATGTCTATCAAA 5 ACAATATCTATTTGAAACAAGACCAGGAGATCCACCTTTTCAAATAACATTTAA TGGAAGAGAAAATATTTAATGGAATCCCAGACTGGGTTTATGAAGAGGA AATGCTTGCTACAAAATATGCTCTCTGGTGGTCTCCTAATGGAAAATTTTTGGCA TATGCGGAATTTAATGATACGGATATACCAGTTATTGCCTATTCCTATTATGGCG ATGAACAATATCCTAGAACAATAAATATTCCATACCCAAAGGCTGGAGCTAAGA 10 ATCCCGTTGTTCGGATATTTATCGATACCACTTACCCTGCGTATGTAGGTCCC CAGGAAGTGCCTGTTCCAGCAATGATAGCCTCAAGTGATTATTATTTCAGTTGGC TCACGTGGGTTACTGATGAACGAGTATGTTTGCAGTGGCTAAAAAGAGTCCAGA ATGTTTCGGTCCTGTCTATATGTGACTTCAGGGAAGACTGGCAGACATGGGATTG TCCAAAGACCCAGGAGCATATAGAAGAAAGCAGAACTGGATGGGCTGGTGGATT 15 CTTTGTTTCAACACCAGTTTTCAGCTATGATGCCATTTCGTACTACAAAATATTTA GTGACAAGGATGGCTACAAACATATTCACTATATCAAAGACACTGTGGAAAATG  ${\tt CTATTCAAATTACAAGTGGCAAGTGGGAGGCCATAAATATATTCAGAGTAACAC}$ AGGATTCACTGTTTTATTCTAGCAATGAATTTGAAGAATACCCTGGAAGAAGAAA 20 CATCTAAGGAAAGAAAGGTGCCAATATTACACAGCAAGTTTCAGCGACTACGCC AAGTACTATGCACTTGTCTGCTACGGCCCAGGCATCCCCATTTCCACCCTTCATG ##ATGGACGCACTGATCAAGAAATTAAAATCCTGGAAGAAAACAAGGAATTGGAAA \*\*ATGCTTTGAAAAATATCCAGCTGCCTAAAGAGGAAATTAAGAAACTTGAAGTAG ATGAAATTACTTTATGGTACAAGATGATTCTTCCTCCTCAATTTGACAGATCAAA 25 GAAGTATCCCTTGCTAATTCAAGTGTATGGTGGTC@CTGCAGTCAGAGTGTAAGG TTGCCTTGGTGGATGGTCGAGGAACAGCTTTCCAAGGTGACAAACTCCTCTATGC AGTGTATCGAAAGCTGGGTGTTTATGAAGTTGAAGACCAGATTACAGCTGTCAG AAAATTCATAGAAATGGGTTTCATTGATGAAAAAAGAATAGCCATATGGGGCTG 30 GTCCTATGGAGGATACGTTTCATCACTGGCCCTTGCATCTGGAACTGGTCTTTTCA CACAGAGAGATTCATGGGTCTCCCAACAAGGATGATAATCTTGAGCACTATAA GAATTCAACTGTGATGGCAAGAGCAGAATATTTCAGAAATGTAGACTATCTTCTC ATCCACGGAACAGCAGATGATAATGTGCACTTTCAGAACTCAGCACAGATTGCT 35 AAAGCTCTGGTTAATGCACAAGTGGATTTCCAGGCAATGTGGTACTCTGACCAGA ACCACGGCTTATCCGGCCTGTCCACGAACCACTTATACACCCACATGACCCACTT CCTAAAGCAGTGTTTCTCTTTGTCAGACTAAAAACGATGCAGATGCAAGCCTGTA **TCAGAATCTGA** 

SEQ ID NO: 684
 zw83d07.s1 Soares\_testis\_NHT Homo sapiens cDNA clone IMAGE:782797 3', mRNA sequence gi|2161864|gb|AA448194.1|AA448194[2161864]
 TTTTTTTTAAAAAAAAAATTAAATATTTTTATTATATACTTTTAAACATATAGAAGA TAGAAAAAAACAGTACAATGAACAGCCATGTCCACCAGTTAGATTCTGTAACAT
 TTTGCCACATACGCCTCACATACATTTTGTTAAACCATTTGAAACATTTTAAGACA CTCTAACACTTCATTCCTAAATGCTTAAGTATGCAAATTAAGACAGTCTTTTATAA ACTACAACACCCTTCTCACAGGCTCATAAAATTACCAATAATTATCCAATATCATT CAAAATCTAATCCACATTCAAATTTTCTCAACTGCCTCACCACCGTGCTGGCCTCC

# CACCCCACCTCAGTCTTTTACAGATGGTTTTTCAAAATAGAGTCCAGTAAAATA TTTCACATTGCATTTGGTTATTACATAACTTT TAATCAAGAAGAGTTAC

**SEQ ID NO: 685** 

- ACTTGCCTCCTCTCTCCCTCTAGAGTCGTGTCTGAACCCGGCCTTTTCCAATTGG CCTGCTCCATCCGAACAGCGTCAACGTGAGTGAATTTGCCCGAAGCTTGTCTTTG CTGAGCGGGTTTGGGGACGTCTGCCCGCCCTCTTTCCCTTCACATTTCATTGCATG GGTTCCCCAACAGCGTTCCCTGGTTCTTTTTTTTGTGACCCCAGTCAATGTCCTGCCT
- 15 CCCCCGGCTCCCGCTCTCTCGCCCCTGGTCTGCGGCGTTCTCTCCGGAATCTTGCC
  CTGGGCCGCGGACGCCCAGGAAAAGAGCCGGGTGCCCCAGGCAGCCTCGCGTTG
  GGGGCGACCGCGCCATCCCGGGAACCGCGAGGCGATCTGAGTCGCCTCCACGTC
  TACCTAAAAGCTGTCGGCCGGGAGGGCGGGCCCCAGAAAGGAGCATTCCTGCG
  GGCTTTTGCTCGACGATCCCCTGCTGAGGCTGTCGCGGGGGGCGAGGGTCCTGCCGAGG

25

**SEQ ID NO: 686** 

yi26g12.s1 Soares placenta Nb2HP Homo sapiens cDNA clone IMAGE:140422 3', mRNA sequence gi|838397|gb|R65759.1|R65759[838397]

- AAAATTTTTTNTACCGTATTTATTGGTTCAAAAACTAGAATTTATAGTTTCAGGCA
  GATTCAACCAAAGAGTCACCAAATTAAATACACAGGGTAGCTTGTGAGGCATA
  GACACAGCCCATGTGTTTTCCTCTACATTGTATATTCATTTCTCTTTTGGCGATTTG
  ACATTATAGCCATTCTCTGGAAGTCCTAAAGCAAACTAGTATTTTATGTGCCATA
  TTAAGTTAAAATTTCTTATGTGAGGATACCACTAATACTGGGTTTTCTATGGATTAAG
  GCATCCTTCTTGCCGGGGGGTATGGACAATGGGGGGTTTTTCTATGGATTAAG
  GNCCCTACCCCTGGGGCCAGGTGNTATGGGGGNATTGTTAAAACCATGGCCATT
- ATTATGGTGGGGGCCAACCCCCACCCNTGGAAG GGGA

SEQ ID NO: 687 >R91550

- 45 AAAGCCTCTGGCCCCACCACATCCCTGTGGGAGAAGATCTGTGAGAAGCTTAAG GAAGAAGGACAGCCAGATATGTGAGCTTAAGTAT GGACAAGCAGATCC

SEQ ID NO: 688 >M94054

GGGCGTGATTTGAGCCCCGTTTTTATTTTCTGTGAGCCACGTCCTCCTCGAGGGG GTCAATCTGGCCAAAAGGAGTGATGCGCTTCGCCTGGACCGTGCTCCTGGG CCTTTGCAGCTCTGCGCGCTAGTGCACTGCGCCCCTCCCGCCGCCGGCCAACAGC AGCCCCGCGCGAGCCGCCGGCGCTCCGGGCGCCTGGCGCCAGCAGATCCAAT 5 GGGAGAACAACGGCAGGTGTTCAGCTTGCTGAGCCTGGGCTCACAGTACCAGC CTCAGCGCCGCGGGACCCGGGCGCCGTCCCTGGTGCAGCCAACGCCTCCG CCCAGCAGCCCGCACTCCGATCCTGCTGATCCGCGACAACCGCACCGCCGCGC GCGAACGCGGACGCCGCTCATCTGGAGTCACCGCTGGCCGCCCCAGGCCCAC CGCCCGTCACTGGTTCCAAGCTGGCTACTCGACATCTAGAGCCCGCGAACGTGGC 10 GCCTCGCGCGGAGAACCAGACAGCGCCGGGAGAAGTTCCTGCGCTCAGTAAC CTGCGGCCGCCAGCCGCGTGGACGGCATGGTGGGCGACGACCCTTACAACCCC TACAAGTACTCTGACGACAACCCTTATTACAACTACTACGATACTTATGAAAGGC CCAGACCTGGGGCAGGTACCGGCCCGGATACGGCACTGGCTACTTCCAGTACG 15 GAAGATGTCCATGTACAACCTGAGATGCGCGGCGGAGGAAAACTGTCTGGCCAG TACAGCATACAGGGCAGATGTCAGAGATTATGATCACAGGGTGCTGCTCAGATTT CCCCAAAGAGTGAAAAACCAAGGGACATCAGATTTCTTACCCAGCCGACCAAGA TATTCCTGGGAATGGCACAGTTGTCATCAACATTACCACAGTATGGATGAGTTTA GCCACTATGACCTGCTTGATGCCAACACCCAGAGGAGAGTGGCTGAAGGCCACA 20 AAGCAAGTTTCTGTCTTGAAGACACATCCTGTGACTATGGCTACCACAGGCGATT TGCATGTACTGCACACACACAGGGATTGAGTCCTGGCTGTTATGATACCTATGGT # GCAGACATAGACTGCCAGTGGATTGATATTACAGATGTAAAACCTGGAAACTAT A SATECTA A AGGTCAGTGTA A ACCCCAGCT A CCTGGTTCCTGA AT CTGACTATACCA MACAATGTTGTGCGCTGTGACATTCGCTACACAGGACATCATGCGTATGCCTCAGG 25 CTGCACAATTTCACCGTATTAGAAGGCAAAGCAAAACTCCCAATGGATAAATCA GTGCCTGGTGTTCTGAAGTGGGAAAAAATAGACTAACTTCAGTAGGATTTATGTA TAACAAAGCACATAACTGGATTTTGAACGCTTAAGTCAATCATTACTTGGAAATT TNTAATGTTTATTATTACATCAACTTTGTGAATTAACACAGTGTTTCAATTCTGT 30 **AATTTCATATTTGACTCTTT** 

**SEQ ID NO: 689** 

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Human mRNA for beta-actin gi|28251|emb|X00351.1|HSAC07[28251] TTGCCGATCCGCCGCCGTCCACACCCGCCGCCAGCTCACCATGGATGATAT CGCCGCGCTCGTCGACAACGCTCCGGCATGTGCAAGGCCGGCTTCGCGGG CGACGATGCCCCCGGGCCGTCTTCCCCTCCATCGTGGGGCGCCCCAGGCACCAG GGCGTGATGGTGGGCATGGGTCAGAAGGATTCCTATGTGGGCGACGAGGCCCAG AGCAAGAGAGGCATCCTCACCCTGAAGTACCCCATCGAGCACGGCATCGTCACC AACTGGGACGACATGGGAAAAATCTGGCACCACACCTTCTACAATGAGCTGCGT GTGGCTCCCGAGGAGCACCCCGTGCTGCTGACCGAGGCCCCCCTGAACCCCAAG GCCAACCGCGAGAGATGACCCAGATCATGTTTGAGACCTTCAACACCCCAGCC ATGTACGTTGCTATCCAGGCTGTGCTATCCCTGTACGCCTCTGGCCGTACCACTG GCATCGTGATGGACTCCGGTGACGGGGTCACCCACACTGTGCCCATCTACGAGG TGACTACCTCATGAAGATCCTCACCGAGCGCGGCTACAGCTTCACCACCACGGCC GAGCGGGAAATCGTGCGTGACATTAAGGAGAAGCTGTGCTACGTCGCCCTGGAC TTCGAGCAAGAGATGGCCACGGCTGCTTCCAGCTCCTCCCTGGAGAAGAGCTAC GAGCTGCCTGACGGCCAGGTCATCACCATTGGCAATGAGCGGTTCCGCTGCCCTG AGGCACTCTTCCAGCCTTCCTTGGGCATGGAGTCCTGTGGCATCCACGAAAC

TACCTTCAACTCCATCATGAAGTGTGACGTGGACATCCGCAAAGACCTGTACGCC AACACAGTGCTGTCTGGCGGCACCACCATGTACCCTGGCATTGCCGACAGGATGC AGAAGGAGATCACTGCCCTGGCACCCAGCACAATGAAGATCAAGATCATTGCTC CTCCTGAGCGCAAGTACTCCGTGTGGATCGGCGGCTCCATCCTGGCCTCGCTGTC 5 CACCTTCCAGCAGATGTGGATCAGCAAGCAGGAGTATGACGAGTCCGGCCCCTC CATCGTCCACCGCAAATGCTTCTAGGCGGACTATGACTTAGTTGCGTTACACCCT TTCTTGACAAAACCTAACTTGCGCAGAAAACAAGATGAGATTGGCATGGCTTTAT 10 TCACAATGTGGCCGAGGACTTTGATTGCACATTGTTGTTTTTTTAATAGTCATTCC AAATATGAGATGCATTGTTACAGGAAGTCCCTTGCCATCCTAAAAGCCACCCCAC TTCTCTCTAAGGAGAATGGCCCAGTCCTCTCCCAAGTCCACACAGGGGAGGTGAT AGCATTGCTTTCGTGTAAATTATGTAATGCAAAATTTTTTTAATCTTCGCCTTAAT ACTTTTTTATTTTGTTTTTTGAATGATGAGCCTTCGTGCCCCCCTTCCCCCTT 15 AGGCAGCCAGGGCTTACCTGTACACTGACTTGAGACCAGTTGAATAAAAGTGCA **CACCTTA** 

**SEQ ID NO: 690** 

20 >AA435938

TTTCATGCTCATTGCTGTTTATTGAAACAAAAGAATCAGAAGAAGATCAGAATGA GAAAGAATTGTAGGAAGGAAAAACTTGTAGAAGTAGAGGGTGGAGAGTGCGAA GATATGTCATGGAAGGCTTCTTTAAACACCCAGAAGAAATTCAGGATAAAGCTCA AAAAGAGCAGGCAATCGATAGGGGTTGAAAATCCACTCAGTAGGCCACGGAAG GACTTCAAGAAGGTTGATCGTTCTGTCGCTGGATGTTGTAGGTGTCCTACGTGAA GGCAATCGACATCTGGATGGCTGTGTGTCTCTTTTGTGTT

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**SEQ ID NO: 691** >AA443497

CGCTGCCTTGCTGGAG

TCCAAGGTCATGGCAAAACATCTGAAGTTCATCGCCAGGACTGTGATGGTACAG GAAGGGAACGTGGAAAGCGCATACAGGACCCTAAACAGAATCCTCACTATGGAT GGGCTCATTGAGGACATTAAGCATCGGCGGTATTATGAGAAGCCATGCCGCCGC 35 GACAGAGGGAAAGCTATGAAAGGTGCCGGCGGATCTACAACATGGAAATGGCTC GCAAGATCAACTTCTTGATGCGAAAGAATCGGGCAGATCCGTGGCAGGGCTGCT GAGGCCTGTGGGTGGGACACCAGTGCGAAACCCTCATCCAGTTTTCTCTCCATCT CTTTTCTTTGTACAATCCCATTTCCTATTACCATTCTCTGCAATAAACTCAAATCA

40 **CATGTCTGC** 

**SEQ ID NO: 692** 

zf17e01.s1 Soares fetal heart NbHH19W Homo sapiens cDNA clone IMAGE:377208 3', mRNA sequence gi|1547536|gb|AA055198.1|AA055198[1547536]

45 CACCTTAAAAACTAGGTTTCTATTTCTGGTTAGATTCTAGAGCAGTGGAACTCAG AGATAACATTGTACAAAACTGTATTTACAAGAAAACCAATTAAAAATTAAGGGT GTGTGCAAAAGTAGACAGGAGAGTCAAGACATATCAATGCAGGGATGGCTTTGG GGAATGGGGACTCAAGGTTCTACACTGGAACCTGGGG

**SEQ ID NO: 693** 

zt87h10.s1 Soares\_testis\_NHT Homo sapiens cDNA clone IMAGE:729379 3', mRNA sequence

- 5 gi|2140847|gb|AA435933.1|AA435933[2140847]
  - TTTTGGTTCAAACAATGGAACATTTTATTATTATCATATTACAAAGAGTCAGTGAT GGGCC
  - ATTCCAGGATTGGTTAATTCAGTAGTTCACCAAAGTCATCAAGGATCCGTCTTTCCATCCCTTCTCTCCCACCCTCAAGGTTTAAGACGGCTGTTGCAGTTCCAGACAT
- 10 TATATCAAGATGCAGTATTCACAGAAAGAGGACTGTTCATTTCTTTACCAGAAGA TTCTCCCATATATCATGTGTCTACATCTAAACCAATCACTACTAAGGGGAAATTG ACCTACAACATTTGGATTAGACTAATCAAATTTACCTTCTGAGTTAGGCATAGAG TCAACTTCTATGAGCACATGGCTGAGCCAAGGATAAGCATTCTGCCAGCAAGAG AGGACATAATATGGGTGTGGGGATTGGAGATGGGAGAG

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- SEQ ID NO: 694
- yo27c07.s1 Soares adult brain N2b5HB55Y Homo sapiens cDNA clone IMAGE:179148 3', mRNA sequence gi|989944|gb|H50103.1|H50103[989944]
- AAATTTATCAATGACAAACAGACATAAAACTCAAAGTTTGGCTCTTCTGAGGGGC
  AGGAGAAAAACTGGTGATGTTCTTTTATACAGATGAAACATGGGTNCAGAAATT
- 20 AGGAGAAAACTGGTGATGTTCTTTTATACAGATGAAACATGGGTNCAGAAATT ACACGNCACTTCTAAAGCAACCAGAAGGGGACACGAAAGCAAACCTGTACATT CAGATTTCCACTAGGTAAGAAAATACANTTTTG
- CAACGTCAGCGTCAGGCGTGTATGAAAAAAAAAAAACCCAGCCGACATGCAG
  - 25 GGGAAGATTTCTCTCATCATTTTTNGTAAANCAAAGCGTTCTAATATTTTACAGA CCAAGTTAGGGCCAGTTTTTNTTTTTCCCT

**SEQ ID NO: 695** 

- za29f01.r1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:293977 5',
- mRNA sequence gi|1267964|gb|N95657.1|N95657[1267964]
  GCAGAAGCGAACAACCTGAGCTTTCCCTTGGAGCCCCTG
  - GCAGAAGCGAACAACCTGAGCTTTCCCTTGGAGCCCTGAGCAGGGAGAGGGCT CACAAGCTTGAGGCCATCTCTCGCCTCTGCGAGNACNAAGTACAAGGACCTAAG AAGATCCGCGAGAAGCGCTCAGCCAGTGCAGACAACCTGACTCTGCCCCGGTGG TCCCCAGCCATCATCTCTTAACTACGGAGGCCCGCCGGACCACACCATCCCTTAG
- 35 TTTCTCCTTTAGTTTGAGAAAAGACAGACTTGGGGTNGGTTTGTTTTTTTC TTTCCTTTTCTTTTTTACGCATAGCTCCCGTCAAAGCTGCCT

**SEQ ID NO: 696** 

- Human lysophosphatidic acid receptor homolog mRNA, complete cds
- 40 gi|1857424|gb|U80811.1|HSU80811[1857424]
  - TCACCACCTACAACCACAGAGCTGTCATGGCTGCCATCTCTACTTCCATCCCTGT AATTTCACAGCCCCAGTTCACAGCCATGAATGAACCACAGTGCTTCTACAACGAG TCCATTGCCTTCTTTTATAACCGAAGTGGAAAGCATCTTGCCACAGAATGGAACA CAGTCAGCAAGCTGGTGATGGGACTTGGAATCACTGTTTGTATCTTCATCATGTT

CGGATGAGCAACCGGCGGGTAGTGGTGGTCATTGTGGTCATCTGGACTATGGCC ATCGTTATGGGTGCTATACCCAGTGTGGGCTGGAACTGTATCTGTGATATTGAAA TTCAACTTGGTGACCTTTGTGGTAATGGTGGTTCTCTATGCTCACATCTTTGGCTA 5 TGTTCGCCAGAGGACTATGAGAATGTCTCGGCATAGTTCTGGACCCCGGCGGAAT CGGGATACCATGATGAGTCTTCTGAAGACTGTGGTCATTGTGCTTGGGGCCTTTA TCATCTGCTGGACTCCTGGATTGGTTTTGTTACTTCTAGACGTGTGCTGTCCACAG TGCGACGTGCTGGCCTATGAGAAATTCTTCCTTCTCCTTGCTGAATTCAACTCTGC CATGAACCCCATCATTACTCCTACCGCGACAAAGAAATGAGCGCCACCTTTAGG 10 CAGATCCTCTGCCAGCGCAGTGAGAACCCCACCGGCCCCACAGAAAGCTCA GACCGCTCGGCTTCCCCCCAACCACCATCTTGGCTGGAGTTCACAGCAATG ACCACTCTGTGGTTTAGAACGGAAACTGAGATGAGGAACCAGCCGTCCTCTTG GAGGATAAACAGCCTCCCCTACCCAATTGCCAGGGCAAGGTGGGGTGTGAGAG AGGAGAAAAGTCAACTCATGTACTTAAACACTAACCAATGACAGTATTTGTTCCT 15 GGACCCCACAAGACTTGATATATTGAAAATTAGCTTATGTGACAACCCTCATC TTGATCCCCATCCCTTCTGAAAGTAGGAAGTTGGAGCTCTTGCAATGGAATTCAA GAACAGACTCTGGAGTGTCCATTTAGACTACACTAACTAGACTTTTAAAAGATTT TGTGTGGTTTGGTGCAAGTCAGAATAAATTCTGGCTAGTTGAATCCACAACTTCA TTTATATACAGGCTTCCCTTTTTTATTTTTAAAGGATACGTTTCACTTAATAAACA 20 CGTTTATGCCTATCAGCAAAAAAAAAAAAAAAAA

SEQ ID NO: 698

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Human interleukin 11 mRNA, complete cds gi|186272|gb|M57765.1|HUMIL11[186272]
GCTCAGGGCACATGCCTCCCCTCCCCAGGCCGGCCCAGCTGACCCTCGGGGCT
CCCCCGGCAGCGGACAGGGAAGGGTTAAAGGCCCCCGGCTCCCTGCCCCTGCC
40 CTGGGGAACCCCTGGCCCTGTGGGGACATGAACTGTGTTTGCCGCCACCACCTGGCC
CCCTCGAGTTTCCCCAGACCCTCGGGCCGAGCTGGACAGCACCACCTGGCC
CCCTCGAGTTTCCCCAGACCCTCGGGCCGAGCTGCACAGCTGAGGGACAAATTC
CCAGCTGACGGGGACCACAACCTGGATTCCCTGCCCACCCTGGCCATGAGTGCG
45 GGGCACTGGGAGCTCTACAGCTCCCAGGTGTGCTGACAAGGCTGCGAGCGGAC
CTACTGTCCTACCTGCGGCACGTGCAGTGCCCGGCAGGTGGCTCTTCCC
TGAAGACCCTGGAGCCCGAGCTGGCACCTGCAGCCCGACCTGGACCGCTGC
TGCGCCGGCTGCAGCTCCTGATGTCCCGCCTGCCCCAGCCACCCCCGGA
CCCGCCGGCGCCCCCCCTGCCCCCTGCCCCAGCCACCCCCGGA

**SEQ ID NO: 699** 

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Homo sapiens mRNA for GABA-BR1a (hGB1a) receptor gi|2826760|emb|Y11044.1|HSGTHLA1[2826760]

15 CGCAGACCCCCAACGCCACCTCAGAAGGTTGCCAGATCATACACCCGCCCTGGG AAGGGGCATCAGGTACCGGGGCCTGACTCGGGACCAGGTGAAGGCTATCAACT TCCTGCCAGTGGACTATGAGATTGAGTATGTGTGCCGGGGGGAGCGCGAGGTGG TGGGGCCCAAGGTCCGCAAGTGCCTGGCCAACGGCTCCTGGACAGATATGGACA 20 CACCCAGCCGCTGTGTCCGAATCTGCTCCAAGTCTTATTTGACCCTGGAAAATGG GAAGGTTTTCCTGACGGGTGGGGACCTCCCAGCTCTGGACGGAGCCCGGGTGGA TTTECGGTGTGACCCCGACTTCCATCTGGTGGGCAGCTCCCGGAGCATCTGTAGT CAGGGCCAGTGGAGCACCCCGAAGCCCCACTGCCAGGTGAATCGAACGCCACAC ##CCAGGGGGCCAGGCCTGCCAGCCGGGGGAGATGGCGCTGGAGGACGTGAAT AGCCGCAGGGACATCCTGCCGGACTATGAGCTCAAGCTCATCCACCACGACAGC AAGTGTGATCCAGGCCAAGCCACCAAGTACCTATATGAGCTGCTCTACAACGAC CCTATCAAGATCATCCTTATGCCTGGCTGCAGCTCTGTCTCCACGCTGGTGGCTG 30

AGGCTGCTAGGATGTGGAACCTCATTGTGCTTTCCTATGGCTCCAGCTCACCAGC

CCTGTCAAACCGGCAGCGTTTCCCCACTTTCTTCCGAACGCACCCATCAGCCACA
CTCCACAACCCTACCCGCGTGAAACTCTTTGAAAAGTGGGGCTGGAAGAAGATT
GCTACCATCCAGCAGACCACTGAGGTCTTCACTTCGACTCTGGACGACCTGGAGG
AACGAGTGAAGGAGGCTGGAATTGAGATTACTTTCCGCCAGAGTTTCTTCTCAGA
TCCAGCTGTGCCCGTCAAAAACCTGAAGCGCCAGGATGCCCGAATCATCGTGGG

35 ACTTTTCTATGAGACTGAAGCCCGGAAAGTTTTTTTGTGAGGTGTACAAGGAGCGT

ACTITICIATGAGACIGAAGCCCGGAAAGITTITITGIGAGGIGIACAAGGAGCGT CTCTTTGGGAAGAAGTACGTCTGGTTCCTCATTGGGTGGTATGCTGACAATTGGT TCAAGATCTACGACCCTTCTATCAACTGCACAGTGGATGAGATGACTGAGGCGGT GGAGGGCCACATCACAACTGAGATTGTCATGCTGAATCCTGCCAATACCCGCAG CATTTCCAACATGACACAGAGAGAAGTTTGTGGAGAAACTAACCAAGCGACTGAA

40 AAGACACCCTGAGGAGACAGGAGGCTTCCAGGAGGCACCGCTGGCCTATGATGC
CATCTGGGCCTTGGCACTGGCCCTGAACAAGACATCTGGAGGAGGCGCCGTTCT
GGTGTGCGCCTGGAGGACTTCAACTACAACAACCAGACCATTACCGACCAAATC
TACCGGGCAATGAACTCTTCGTCCTTTGAGGGTGTCTCTGGCCATGTGGTGTTTG
ATGCCAGCGGCTCTCGGATGGCATGACCAAGGATGATCTTTCCTGGTCCAA

45 GCTACAAGAAGATTGGCTACTATGACAGCACCAAGGATGATCTTTCCTGGTCCAA
AACAGATAAATGGATTGGAGGGTCCCCCCCAGCTGACCAGACCCTGGTCATCAA
GACATTCCGCTTCCTGTCACAGAAACTCTTTATCTCCGTCTCAGTTCTCCAGCC
TGGGCATTGTCCTAGCTGTTGTCTGTCTGTCCTTTAACATCTACAACTCACATGTC
CGTTATATCCAGAACTCACAGCCCAACCTGAACAACCTGACTGCTGTGGGCTGCT

CACTGGCTTTAGCTGCTGTCTTCCCCCTGGGGCTCGATGGTTACCACATTGGGAG GAACCAGTTTCCTTTCGTCTGCCAGGCNCGCCTCTGGCTCCTGGGCCTGGGCTTTA GTCTGGGCTACGGTTCCATGTTCACCAAGATTTGGTGGGTCCACACGGGCTTCAC AAAGAAGGAAGAAAGAAGGAGTGGAGGAAGACTCTGGAACCCTGGAAGCTGT 5 ATGCCACAGTGGGCCTGCTGGTGGGCATGTCCTCACTCTCGCCATCTGGCA GATCGTGGACCCTCTGCACCGGACCATTGAGACATTTGCCAAGGAGGAACCTAA GGAAGATATTGACGTCTCTATTCTGCCCCAGCTGGAGCATTGCAGCTCCAGGAAG ATGAATACATGGCTTGGCATTTTCTATGGTTACAAGGGGCTGCTGCTGCTGCTGG GAATCTTCCTTGCTTATGAGACCAAGAGTGTGTCCACTGAGAAGATCAATGATCA 10 CCGGGCTGTGGCATGGCTATCTACAATGTGGCAGTCCTGTGCCTCATCACTGCT CCTGTCACCATGATTCTGTCCAGCCAGCAGGATGCAGCCTTTGCCTTTGTCTCT TGCCATAGTTTTCTCCTCCTATATCACTCTTGTTGTGCTCTTTTGTGCCCAAGATGC GCAGGCTGATCACCCGAGGGGAATGGCAGTCGGAGGCGCAGGACACCATGAAG ACAGGGTCATCGACCAACAACAACGAGGAGGAGAAGTCCCGGCTGTTGGAGAA 15 GGAGAACCGTGAACTGGAAAAGATCATTGCTGAGAAAGAGGAGCGTGTCTCTGA ACTGCGCCATCAACTCCAGTCTCGGCAGCAGCTCCGGTCCCGGCGCCACCACCG ACACCCCAGAACCCTCTGGGGGCCTGCCCAGGGGACCCCCTGAGCCCCCGAC CGGCTTAGCTGTGATGGGAGTCGAGTGCATTTGCTTTATAAGTGAGGGTAGGGTG AGGGAGGACAGGCCAGTAGGGGGAGGGGAAAGGGAGAGGGGAAGGGCAGGGGA 20 CTCAGGAAGCAGGGGTCCCCATCCCAGCTGGGAAGAACATGCTATCCAATCT CATCTCTTGTAAATACATGTCCCCCTGTGAGTTCTGGGCTGATTTGGGTCTCTCAT ACGGCAACCCCTGCAGCTCCTCTGCCTTTGTGCTCTGTTCCTGTCCAGCAGGGGTC 25 TCCCAACAAGTGCTCTTTCCACCCCAAAGGGGCCTCTCCTTTTCTCCACTGTCATA ATCTCTTTCCATCTTACTTGCCCTTCTATACTTTCTCACATGTGGCTCCCCCTGAAT TTTGCTTCCTTTGGGAGCTCATTCTTTTCGCCAAGGCTCACATGCTCCTTGCCTCT GCTCTGTGCACTCACGCTCAGCACACATGCATCCTCCCCTCTCCTGCGTGTGCCCA 30 CTGAACATGCTCATGTTACACACGCTTTTCCCGTATGCTTTCTTCATGTTCAGTC ACATGTGCTCTCGGGTGCCCTGCATTCACAGCTACGTGTGCCCCTCTCATGGTCAT GGGTCTGCCCTTGAGCGTGTTTGGGTAGGCATGTGCAATTTGTCTAGCATGCTGA GTCATGTCTTTCCTATTTGCACACGTCCATGTTTATCCATGTACTTTCCCTGTGTAC CCTCCATGTACCTTGTGTACTTTCTTCCCTTAAATCATGGTATTCTTCTGACAGAG 35 CCATATGTACCCTACCCTGCACATTGTTATGCACTTTTCCCCAATTCATGTTTGGT GGGGCCATCCACACCCTCTCCTTGTCACAGAATCTCCATTTCTGCTCAGATTCCCC CCATCTCCATTGCATTCATGTACTACCCTCAGTCTACACTCACAATCATCTTCTCC CAAGACTGCTCCCTTTTGTTTTTGTGTTTTTTTGAGGGAAATTAAGGAAAAATAAG TGGGGCAGGTTTGGAGAGCTGCTTCCAGTGGATAGTTGATGAGAATCCTGACC 40 AAAGGAAGCACCCTTGACTGTTGGGATAGACAGATGGACCTATGGGGTGGGAG GTGGTGTCCCTTTCACACTGTGGTGTCTCTTGGGGAAGGATCTCCCCGAATCTCA 

**SEQ ID NO: 700** 

zh96g08.s1 Soares\_fetal\_liver\_spleen\_1NFLS\_S1 Homo sapiens cDNA clone IMAGE:429182 3', mRNA sequence gi|1448327|gb|AA004759.1|AA004759[1448327] ACTTTATGCAAAAAAAAAAATATACATTTATTTATAGGTCTCAATACAGCAAAATGA AAACGAAAATTGAGAACATTGCTCATTAGGCCAGCAACTTTAAAATTATTTAATT TGAAATATAAAATAGGTGGTCTTCATAAAAAAGATGCATGAAATTACCTTACCTT

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**SEQ ID NO: 701** 

Homo sapiens canalicular multispecific organic anion transporter 2 (CMOAT2) mRNA, complete cds gi|3550323|gb|AF083552.1|AF083552[3550323]

AGCCGCGCCTCGGCCCCATGGACGCCCTGTGCGGTTCCGGGGAGCTCGGCTCCAA

10 GTTCTGGGACTCCAACCTGTCTGTGCACACAGAAAACCCGGACCTCACTCCCTGC
 TTCCAGAACTCCCTGCTGGCCTGGGTGCCCTGCATCTACCTGTGGGTCGCCCTGC
 CCTGCTACTTGCTCTACCTGCGGCACCATTGTCGTGGCTACATCATCCTCTCCAC
 CTGTCCAAGCTCAAGATGGTCCTGGGTGTCCTGTGGTGCGTCTCCTGGGCGG
 ACCTTTTTACTCCTTCCATGGCCTGGTCCATGGCCGGGCCCCTGCCCCTGTTTTC

15 TTTGTCACCCCCTTGGTGGTGGGGGTCACCATGCTGCTGCCACCTGCTGATAC
 AGTATGAGCGGCTGCAGGGCGTACAGTCTTCGGGGGGTCCTCATTATCTTCTGGTT
 CCTGTGTGTGGTGTCCCATCGTCCCATTCCGCTCCAAGATCCTTTTAGCCAAGG
 CAGAGGGTGAGATCTCAGACCCCTTCCGCTTCACCACCTTCTACATCCACTTTGC

30 GTTTCGTACTGGGATCATGGGTGTCATCTACAGGAAGGCTCTGGTTATCACCAAC
 TCAGTCAAACGTGCGTCCACTGTGGGGGAAATTGTCAACCTCATGTCAGTGGATG
 CCCAGCGCTTCATGGACCTTGCCCCCTTCCTCAATCTGCTGTGGTCAGCACCCCTG
 CAGATCATCCTGGCGATCTACTTCCTCTGGCAGAACCTAGGTCCCTCTGTCCTGG
 CTGGAGTCGCTTTCATGGTCTTGCTGATTCCACTCAACGGAGCTGTGGCCGTGAA

 35 GATGCGCGCCTTCCAGGTAAAGCAAATGAAATTGAAGGACTCGCGCATCAAGCT
 GATGAGTGAGATCCTGAACGGCATCAAGGTGCTGAAGCTGTACGCCTGGGAGCC

CAGCTTCCTGAAGCAGGTGGAGGCATCAGGCAGGTGAGCTCCAGCTGCTGCGCACGGCGGCCTACCTCCACACCACCACCACCACCTTCACCTGGATGTGCAGCCCCTTC

TTGGTGACCCTGATCACCCTCTGGGTGTACGTGTACGTGGACCCAAACAATGTGC

TGGACGCCGAGAAGGCCTTTGTGTCTGTGTCCTTGTTTAATATCTTAAGACTTCCC
CTCAACATGCTGCCCCAGTTAATCAGCAACCTGACTCAGGCCAGTGTGTCTCTGA
AACGGATCCAGCAATTCCTGAGCCAAGAGGAACTTGACCCCCAGAGTGTGGAAA
GAAAGACCATCTCCCCAGGCTATGCCATCACCATACACAGTGGCACCTTCACCTG
GGCCCAGGACCTGCCCCCCACTCTGCACAGCCTAGACATCCAGGTCCCGAAAGG

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GAGAGAAGGCATTAACCTGTCTGGGGGCCAGCGGCAGCGGGTCAGTCTGGCTC GAGCTGTTTACAGTGATGCCGATATTTTCTTGCTGGATGACCCACTGTCCGCGGT GGACTCTCATGTGGCCAAGCACATCTTTGACCACGTCATCGGGCCAGAAGGCGTG CTGGCAGGCAAGACGCGAGTGCTGGTGACGCACGGCATTAGCTTCCTGCCCCAG ACAGACTTCATCATTGTGCTAGCTGATGGACAGGTGTCTGAGATGGGCCCGTACC CAGCCTGCTGCAGCGCAACGGCTCCTTTGCCAACTTTCTCTGCAACTATGCCCC CGATGAGGACCAAGGGCACCTGGAGGACAGCTGGACCGCGTTGGAAGGTGCAG AGGATAAGGAGCACTGCTGATTGAAGACACACTCAGCAACCACACGGATCTGA CAGACAATGATCCAGTCACCTATGTGGTCCAGAAGCAGTTTATGAGACAGCTGA GTGCCTGTCCTCAGATGGGGAGGGACAGGGTCGGCCTGTACCCCGGAGGCACC TGGGTCCATCAGAGAAGGTGCAGGTGACAGAGGCGAAGGCAGATGGGGCACTG ACCCAGGAGGAGAAAGCAGCCATTGGCACTGTGGAGCTCAGTGTTCTGGGAT TATGCCAAGGCCGTGGGGCTCTGTACCACGCTGGCCATCTGTCTCCTGTATGTGG GTCAAAGTGCGGCTGCCATTGGAGCCAATGTGTGGCTCAGTGCCTGGACAAATG ATGCCATGGCAGACAGTAGACAGAACACTTCCCTGAGGCTGGGCGTCTATG CTGCTTTAGGAATTCTGCAAGGGTTCTTGGTGATGCTGGCAGCCATGGCCATGGC AGCGGGTGGCATCCAGGCTGCCCGTGTGTTGCACCAGGCACTGCTGCACAACAA GATACGCTCGCCACAGTCCTTCTTTGACACCACCACCATCAGGCCGCATCCTGAAC TGCTTCTCCAAGGACATCTATGTCGTTGATGAGGTTCTGGCCCCTGTCATCCTCAT GCTGCTCAATTCCTTCTAACGCCATCTCCACTCTTGTGGTCATCATGGCCAGCA CGCCGCTCTTCACTGTGGTCATCCTGCCCCTGGCTGTGCTCTACACCTTAGTGCAG A CONTRACTOR OF THE PROPERTY O \*\*\*\*\*\*\*\*CTACAACCGCAGCCGGATTTTGAGATCATCAGTGATACTAAGGTGGATGCCAA \*\*\* CCAGAGAAGCTGCTACCCCTACATCTCCCAACCGGTGGCTGAGCATCGGAGTG GAGTTCGTGGGAACTGCGTGGTGCTCTTTGCTGCACTATTTGCCGTCATCGGGA GGAGCAGCCTGAACCCGGGGCTGGTGGGCCTTTCTGTGTCCTACTCCTTGCAGGT GACATTTGCTCTGAACTGGATGATACGAATGATGTCAGATTTGGAATCTAACATC GTGGCTGTGGAGAGGGTCAAGGAGTACTCCAAGACAGAGACAGAGGCGCCCTGG GTGGTGGAAGGCAGCCCCCCCGAAGGTTGGCCCCCACGTGGGGAGGTGGAG TTCCGGAATTATTCTGTGCGCTACCGGCCGGGCCTAGACCTGGTGCTGAGAGACC TGAGTCTGCATGTGCACGGTGGCGAGAAGGTGGGGATCGTGGGCCGCACTGGGG CTGGCAAGTCTTCCATGACCCTTTGCCTGTTCCGCATCCTGGAGGCGGCAAAGGG TGAAATCCGCATTGATGGCCTCAATGTGGCAGACATCGGCCTCCATGACGTGCGC TCTCAGCTGACCATCATCCCGCAGGACCCCATCCTGTTCTCGGGGACCCTGCGCA TGAACCTGGACCCCTTCGGCAGCTACTCAGAGGAGGACATTTGGTGGGCTTTGGA GCTGTCCCACCTGCACACGTTTGTGAGCTCCCAGCCGGCAGGCCTGGACTTCCAG TGCTCAGAGGGCGGGAGAATCTCAGCGTGGGCCAGAGGCAGCTCGTGTGCCTG GCCCGAGCCTGCTCCGCAAGAGCCGCATCCTGGTTTTAGACGAGGCCACAGCTG CCATCGACCTGGAGACTGACAACCTCATCCAGGCTACCATCCGCACCCAGTTTGA TACCTGCACTGTCCTGACCATCGCACACCGGCTTAACACTATCATGGACTACACC AGGGTCCTGGTCCTGGACAAAGGAGTAGTAGCTGAATTTGATTCTCCAGCCAACC TCATTGCAGCTAGAGGCATCTTCTACGGGATGGCCAGAGATGCTGGACTTGCCTA GACACCAAATATGTCCGCAGAATGGACTTGATAGCAAACACTGGGGGCACCTTA AGATTTTGCACCTGTAAAGTGCCTTACAGGGTAACTGTGCTGAATGCTTTAGATG AGGAAATGATCCCCAAGTGGTGAATGACACGCCTAAGGTCACAGCTAGTTTGAG CCAGTTAGACTAGTCCCCGGTCTCCCGATTCCCAACTGAGTGTTATTTGCACACT 

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SEQ ID NO: 702

yq42d10.s1 Soares fetal liver spleen 1 NFLS Homo sapiens cDNA clone IMAGE:198451 3', mRNA sequence gi|970054|gb|R94659.1|R94659[970054]

TTGTTTTTTTGGTTCAGCATAACTTGGAACATTTGAAAGCTTTTCAACCTAAATG

10 TGGG

GAAAAAACAGGTAAGGCATTATTTTTGCACAAAACTAGCATTCCTAATAGTGCA AATGAA

 ${\tt TCTGATACCTCTTAAAAATGGTGAGAGGTCATACACTTACTAGATTAGATT}\\ {\tt TTCTT}$ 

15 TCTATGGCTTGACAAATTATCCCTCTATAAATTCTACTCTCACCCAGAGGCTGTTG CTGT

AATCAAAAGGATAACTGTAGGATAAAGGTCCAACCTTCTCCTGGTATCCGGCAA AAGGGT

TTTTGCTCATATGGCAAAAAAAATCTAATTTTTAAATTATCCTACAGNGGAATAT

20 ACAAC

TGGGNTTCCTNGGGACCCTCTATTTATCNGGCGGCAACAGGTGGTTCGGGGCGCCCGC
GGGGGCCCCTAACCCAAAATTGGGCGGCAAATCT

TO SEED ID NOT 703 WE TO THE SEED RESERVED AND A CONTROL OF THE SEED RESERVED TO A CO

25 zd29f03.s1 Soares\_fetal\_heart\_NbHH19W Homo sapiens cDNA clone IMAGE:342077 3', mRNA sequence gi|1367074|gb|W60315.1|W60315[1367074] CATAACTTAAGTAAACTTTATTTTCAAAATGCTTCAGGTACAAAAGAAAACAATC GGCAAAGTCTAACAATAATTAACAAACCAGCTCTTGAGCGGCAGAGTGCTCCAG GGATGAGAGGGGCTGGGGATGGAAAGGTGGTTGGGAGACACAACATTTTTCTAG

DESTRUCTION OF THE CONTROL OF THE CONTROL OF SAME OF THE PROPERTY OF THE SAME OF THE SAME

30 CTTCAGAAAGTCAGGGAGCCCAGATCACAGCCTGAACTTCATGGTATTGGTTACA GATTCTTTACAAAGGTGTTTACCTCTCTCATGAGGTCTTCTTGATTGGTTACTTCC TCAGAAAAATCATCATTGACATCCAACACCAGCACTGGAATGTTCATCAGAGCCT CAAAGTGGAGCCTGTCACTTGTACACANGACCTCTCAAAGATCTGTACTGGCTTC CTGGCCTGGTAAGAGTTCTCAGGGGAAG

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**SEQ ID NO: 704** 

yb54f05.r1 Stratagene ovary (#937217) Homo sapiens cDNA clone IMAGE:75009 5', mRNA sequence gi|653755|gb|T51895.1|T51895[653755]

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**SEQ ID NO: 705** 

zx69a01.s1 Soares\_total\_fetus\_Nb2HF8\_9w Homo sapiens cDNA clone IMAGE:796680 3', mRNA sequence gi|2185799|gb|AA460679.1|AA460679[2185799] TACTCAGTCACCACCCAGAAATTGTCCGAGTTATGAAAATAGATTCATTTTGAGAA

# **SEQ ID NO: 706**

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- 15 AAAGTCAACAGAACCTCCACTAGGCATAATTTACATATGTACAGACTCAATCAGC TTTTAATATAGAAAGATATTTGAACCCAAAATCTTTCATTAAGGTAAAAAATACA ATAATAATTTTAATGAAATCCTGGAAAAATTCATACAAATAAAATTAAAAGCCTC CAATGGGGTATAATCCAGCAATATCCTAGGCAAATGCCTCCTGAAGAACAACAG CCTTTTTAAAAACATCACTGTTTATCATTCAAAAATTCAGACGTCTCCTATCTTTGGC
- 20 TATTTTATCTCTTCAACT

#### - 1.074/ASEO ID INO: 707 (2.13) (4.16) (4.16) (4.16) (4.17) (4.17) (4.17) (4.17) (4.17) (4.17) (4.17) (4.17)

aa47b01.r1 NCI\_CGAP\_GCB1 Homo sapiens cDNA clone TMAGE:824041 5! similar to TR:G1049078 G1049078 SRP30C.;, mRNA sequence

- 25 gi|2219894|gb|AA490721.1|AA490721[2219894]
  TATCTCAGAAAAGAAGACATGCGATATGCCCTGCGTAAACTGGATGACACCAAA
  TTCCGCTCTCATGAGGGTGAAACTTCCTACATCCGAGTTTATCCTGAGAGAAGCA
  CCAGCTATGGCTACTCACGGTCTCGGTCTAGGGCCCGTGACTCTCCATA
  CCAAAGCAGGGGTTCCCACACTACTTCTCTCTTCAGGCCCTACTGAGACAGGT

## SEQ ID NO: 708

- Human 78 kdalton glucose-regulated protein (GRP78) gene, complete cds
  gi|183644|gb|M19645.1|HUMGRP78[183644]
  CCCGGGGTCACTCCTGCTGGACCTACTCCGACCCCCTAGGCCGGAGTGAAGGC
  GGGACTTGTGCGGTTACCAGCGGAAATGCCTCGGGGTCAGAAGTCGCAGGAGAG
  ATAGACAGCTGCTGAACCAATGGGACCAGCGGATGGGGCGGATGTTATCTACCA

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ACCTACTCCTGGTAAGTGGGGTTGCGGATGAGGGGGACGGGGCGTGGCGCTGGC TGGCGTGAGAAGTGCGGTGCTGATGTCCCTCTGTCGGGTTTTTGCAGCGTCGGCG TGTTCAAGAACGCCGCGTGGAGATCATCGCCAACGATCAGGGCAACCGCATCA CGCCGTCCTATGTCGCCTTCACTCCTGAAGGGGAACGTCTGATTGGCGATGCCGC CAAGAACCAGCTCACCTCCAACCCGAGAACACGGTCTTTGACGCCAAGCGGCT CATCGGCCGCACGTGGAATGACCCGTCTGTGCAGCAGGACATCAAGTTCTTGCCG TATTTAGAGTTATAAGTCTCTGGAAAAGTGTTGAGACAACAGTTGAAGGTTATAG ACATGATGTATGTAATAACTTTAATACTATTAGTATGTTACAAAACTTAAGACAG 10 TTGCTGTCGTACTGTCTACGATAGTTTAGGAATAAAAGACCGATTAAAACTGAAC TTTGTAAGACACCTATACTCCCTGAAGTATTTCTAGTCAATTTGCAGCCCCAAGG GACCAAAATAAACCAAATTGTGGGGATGGTAGTGGGTCTTTTAAACTTTGAGATG TCATTGTATCTGTGTCTGAAAACAATAATTCTTTAAAATAGGTGGTTGAAAAGAA TATTTGGGAAAGAAGGTAAATATTTCTAGAACAATGTTAAGTATTTTTTGATCAT TAGTATTCTCGGTTGGCTGTTATGTATAGAAGCCTTCGTGAAGGGTTTCAAAAAT TTTAATCAGAATGGTATTCATGCTTGTCACGGTTTAATTATTGAGTCCCTTTACTA TAAGCCAAACAAAATAGACTTTTCATGTATTATTTAATGCTTACAATTCCAGGA 20 ACAATAAAATTTTATATGTTGTATTCATCAATAATTGGCTTAAAAAACTAAAGTGA TGGTTTGACTGTAATTTTTTTTTTTGAGATGGAGTCTTGCTCTGTTGCCCAGGCT GACTGCAGTGCACGATCTCAGCTCACTGCAACCTCTGCCTCCCGGGTTAAGCA GCTCTCCTGCCTCAGCCTCAAGTAATGGAACGACAGGCACACCACCACAGGTG GCTAATTTTTTTTTTTTTTTAATTTTCAGTAGAGACAGGGTTTCTCCACATTGCC AGGCTGGTCTTGAAATCCTGCCCTCAGGTTGATCCTCCTGCCTAGCCTCCCAAAG TGCTGGATTATAGGCAGAAGCCACCGCCTGGCCAGACTGTAATTTAAATAAGGG TTAAACTATGTGACAATACACTTAATTATCTTTATCCTTTTAGGTTACCCATGCAG TTGTTACTGTACCAGCCTATTTTAATGATGCCCAACGCCAAGCAACCAAAGACGC TGGAACTATTGCTGGCCTAAATGTTATGAGGATCATCAACGAGCCGTAAGTATGA 30 AATTCAGGGATACGGCATATTTGCCAAATAGTGGAAATGTGAAGTACTGACAAA ACTTTTCCCTTTTTCAATCTAATAGTACGCCAGCTGCTATTGCTTATGGCCTGGAT AAGAGGGAGGGGAGAAGAACATCCTGGTGTTTGACCTGGGTGGCGGAACCTTC GATGTGTCTCTCACCATTGACAATGGTGTCTTCGAAGTTGTGGCCACTAATG GAGATACTCATCTGGGTGGAGAAGACTTTGACCAGCGTGTCATGGAACACTTCAT 35 TGCAGAAACTCCGGCGCGAGGTAGAAAAGGCCAAGGCCCTGTCTTCTCAGCATC AAGCAAGAATTGAAATTGAGTCCTTCTATGAAGGAGAAGACTTTTCTGAGACCCT GACTCGGGCCAAATTTGAAGAGCTCAACATGGTATGTTCCTTGTTTTCTGCTTTGC TAATGAGATCTCCTTAGACTCTGAATTCAGGACATTGCATCTAGATACTTAGATA 40 ACAGACATCACAGTAACCATGTCTTTTTTCTAGGATCTGTTCCGGTCTACTATGAA GCCCGTCCAGAAAGTGTTGGAAGAATTCTGATTTGAAGAAGTCTGATATTGATGAA ATTGTTCTTGTTGGTGGCTCGACTCGAATTCCAAAGATTCAGCAACTGGTTAAAG AGTTCTTCAATGGCAAGGAACCATCCCGTGGCATAAACCCAGATGAAGCTGTAG CGTATGGTGCTGCTCCAGGCTGGTGTGCTCTCTGGTGATCAAGATACAGGTAG GTCATCATCGCAGCATCTTTCTTAGTGATTCAGTAGCTTGATGGAAGAGCTCGGT ACCCCTATTGCTTTAGAAAATACCAGAATATGAGCAACAAGGTCACACAGCTAG TAAAGGGTATAAGTGAAGACAAGACTGGGGTAGTCTCCAAGATCATTAGCAACT GTTTAATTCACTGCCTTTAAAATGTGTGTGTTAGAACCTAACCAAATGTTAGAGA GATAAACTTTACATAGCTCATAGGGAGAACTTGAATTAAAAGTTAAATAACTTAT

CCTTACAGGTGACCTGGTACTGCTTCATGTATGTCCCCTTACACTTGGTATTGAAA CTGTAGGAGGTGTCATGACCAAACTGATTCCAAGTAATACAGTGGTGCCTACCAA 5 GAAGTCTTGCTCTGTTGCCCAGGCTGGACTGCAGTGGCACGATCTCGGCTCACTG CAAATTCTGTCTCCCGGGTTCAAGTGATTCTCCTGCCTCAGCCTCCAGAGTAGCT GGATTACAGCCTGACCACCACACCTGGCTAATTTCTGTATTTTTAGTAGAGGATG GGCTTTCACCATGTTTCCCAGGCTGGTCTCCAACTCCTGACCTCAGGTCATCTGCC TGCCTCCACCGTCCCGAAAGTACTGGGATTATAGCGTGAGCCACCACGCCAGATC 10 TATCTATCATGGCATATTTTAAAAGAACATGACTTAATATGTCCTATTGAAATGG CTAGGGAACTAAGTAACTGCTGTTTTCAGATGGAGGTCTTAATTTGAATAATGTT GATATTAGATATTTAGCATTCTTTTTTTTTTTTTTAATGGAGTCTTGCTCTGTCG CCTAGGCTGGGTGCAGTGGCATGACTTGCAACCTCTGCCTCCCGAATAGCTGGG ATTACAGGTGCCCACCATCACGCCCGGCTAAGTTTTGTATTTTTAGTAGAGGCGA 15 GTTTCGCCATGTTGGCCAGGCTGGTCTTGAACCCCTAACCTCAGTGATCCCACGG TCACCGACCTGGCCTCCCAAAAGTACTGTACCCAGCCAATGATTAGCATTCTCAC TAATAATAGCATCTGAGCTGGCTCCTAGAGTACAAGAAAAAGGAGTTCACAGTA CTTTAAAATAGATAAAATTCAGTTGAGTTAGTAACCTAACTCATTGTTAGTACTA GTTGCTGCTCCTTGTAGACCAATATGAAATTACTTTTAGCTCGATAAAACCAAAA 20 GTGTCACTTTATGCTTCAGACTGAAATGCGGGGATCTAGATGTGCTAATGCTTGT CAGTAACAACTAACAAGTTTTTCTGTATGTAACTTCTAGGTGAAAGACCCCTGAC MAAAAGACAATCATCTTCTGGGTACATTTGATCTGACTGGAATTCCTCCTGCTCCTC Maria de GTGGGGTCCCACAGATTGAAGTCACCTTTGAGATAGATGTGAATGGTATTCTTCG 😘 AGTGACAGCTGAAGACAAGGGTACAGGGAACAAAAATAAGATCACAATCACCA 25 ATGACCAGAATCGCCTGACACCTGAAGAAATCGAAAGGATGGTTAATGATGCTG AGAAGTTTGCTGAGGAAGACAAAAAGCTGAAGGAGCGCATTGATACTAGAAATG AGTTGGAAAGCTATGCCTATTCTCTAAAGAATCAGATTGGAGATAAAGAAAAGC TGGGAGGTAAACTTTCCTCTGAAGATAAGGAGACCATGGAAAAAGCTGTAGAAG AAAAGATTGAATGGCTGGAAAGCCACCAAGATGCTGACATTGAAGACTTCAAAG 30 CTAAGAAGAAGGAACTGGAAGAAATTGTTCAACCAATTATCAGCAAACTCTATG GAAGTGCAGGCCCTCCCCCAACTGGTGAAGAGGATACAGCAGAAAAAGATGAGT TGTAGACACTGATCTGCTAGTGCTGTAATATTGTAAATACTGGACTCAGGAACTT TTGTTAGGAAAAATTGAAAGAACTTAAGTCTCGAATGTAATTGGAATCTTCACC TCAGAGTGGAGTTGAAACTGCTATAGCCTAAGCGGCTGTTTACTGCTTTTCATTA 35 GCAGTTGCTCACATGTCTTTGGGTGGGGGGGAGAAGAAGAATTGGCCATCTTAA AAAGCGGGTAAAAACCTGGGTTAGGGTGTGTGTCACCTTCAAAATGTTCTATT TAACAACTGGGTCATGTGCATCTGGTGTAGGAGGTTTTTTCTACCATAAGTGACA CCAATAAATGTTTGTTATTTACACTGGTCTAATGTTTGTGAGAAGCTT

40 SEO ID NO: 709

ACCAACTACTTGTGGTGTCACTGGCGGCGGCCGACATCGCAGTGGGTGTGCTCG CCTCTTCATTGCCTGCTTCGTCCTGGTCCTCACGCAGAGCTCCATCTTCAGTCTCC TGGCCATCGCCATTGACCGCTACATTGCCATCCGCATCCCGCTCCGGTACAATGG 5 CTTGGTGACCGGCACGAGGGCTAAGGGCATCATTGCCATCTGCTGGGTGCTGTCG TTTGCCATCGCCTGACTCCCATGCTAGGTTGGAACAACTGCGGTCAGCCAAAGG AGGGCAAGAACCACTCCCAGGGCTGCGGGGAGGGCCAAGTGGCCTGTCTCTTTG AGGATGTGGTCCCCATGAACTACATGGTGTACTTCAACTTCTTTGCCTGTGTGCTG GTGCCCTGCTGCTCATGCTGGGTGTCTATTTGCGGATCTTCCTGGCGGCGCGAC 10 GACAGCTGAAGCAGATGGAGAGCCAGCCTCTGCCGGGGGAGCGGGCACGGTCCA CACTGCAGAAGGAGGTCCATGCTGCCAAGTCACTGGCCATCATTGTGGGGCTCTT TGCCCTCTGCTGGCTGCCCCTACACATCATCAACTGCTTCACTTTCTTCTGCCCCG ACTGCAGCCACGCCCTCTCTGGCTCATGTACCTGGCCATCGTCCTCTCCCACACC AATTCGGTTGTGAATCCCTTCATCTACGCCTACCGTATCCGCGAGTTCCGCCAGA 15 CCTTCCGCAAGATCATTCGCAGCCACGTCCTGAGGCAGCAAGAACCTTTCAAGGC AGCTGGCACCAGTGCCCGGGTCTTGGCAGCTCATGGCAGTGACGGAGAGCAGGT CAGCCTCCGTCTCAACGCCACCCGCCAGGAGTGTGGGCCAACGGCAGTGCTCC TGCCCAAGAGTCCCAGGGGAACACGGGCCTCCCAGACGTGGAGCTCCTTAGCCA 20 TGAGCTCAAGGGAGTGTGCCCAGAGCCCCCTGGCCTAGATGACCCCCTGGCCCA GGATGGAGCAGGAGTGTCCTGATGATTCATGGAGTTTGCCCCTTCCTAAGGGAAG PARTICIPAGE AGAGE CONTROL OF THE STATE OF TH 25 'AGATGTTTCATGCTGTGAGGCCTTGCACCAGGTGGGGGCCACAGCACCAGCAGC ATCTTTGCTGGGCAGGGCCCAGCCCTCCACTGCAGAAGCATCTGGAAGCACCACC TTGTCTCCACAGAGCAGCTTGGGCACAGCAGACTGGCCTGGGCCCTGAGACTGGG GAGTGGCTCCAACAGCCTCCTGCCACCCACACCACTCTCCCTAGACTCTCCTA GGGTTCAGGAGCTGCTGGGCCCAGAGGTGACATTTGACTTTTTTCCAGGAAAAAT GTAAGTGTGAGGAAACCCTTTTTATTTATTACCTTTCACTCTCTGGCTGCTGGGT 30 CTGCCGTCGGTCCTGCTAACCTGGCACCAGAGCCTCTGCCGGGGAGCCTCAG GCAGTCCTCTCCTGCTGTCACAGCTGCCATCCACTTCTCAGTCCCAGGGCCATCTC TTGGAGTGACAAAGCTGGGATCAAGGACAGGAGTTGTAACAGAGCAGTGCCAG AGCATGGGCCCAGGTCCCAGGGGAGAGGTTGGGGCTGGCAGGCCACTGGCATGT GCTGAGTAGCGCAGAGCTACCCAGTGAGAGGCCTTGTCTAACTGCCTTTCCTTCT 35 AAAGGGAATGTTTTTTCTGAGATAAAATAAAACGAGCCACATCGTGTTTTAAG CTTGTCCAAATGAAAAAAAAAAAAAAAAAAAA

#### SEQ ID NO: 710

NAAGCCTGGTAAGAATTGGGGGGAACCCACTTGGTATTGNCCCTCTTCCAGGATT TTGGAAATTCCAACCGGCCTTGGNTTTAAGAGAAAANAAGGGNTGGTTCCCACT AAT

- 5 SEQ ID NO: 711
  - ab36c08.r1 Stratagene HeLa cell s3 937216 Homo sapiens cDNA clone IMAGE:842894 5' similar to TR:G1256802 G1256802 SODIUM/POTASSIUM-TRANSPORTING ATPASE BETA-3 SUBUNIT.; mRNA sequence gi|2218877|gb|AA489275.1|AA489275[2218877] CTGGCCGAGTGGAAGCTCTTCATCTACAACCCGACCACCGGAGAATTCCTGGGGC
- 10 GCACCGCAAGAGCTGGGGTTTGATCTTGCTCTTCTACCTAGTTTTTTATGGGTTCC
  TGGCTGCACTCTTCTCATTCACGATGTGGGTTATGCTTCAGACTCTCAACGATGA
  GGTTCCAAAAATACCGTGACCAGATTCCTAGCCCAGGACTCATGGTTTTTCCAAAA
  CCAGTGACCGCATTGGAATATACATTCAGTAGGTCTGATCCAACTTCGTATGCAG
  GGTACATTGAAGACCTTAAGAAGTTTCTAAAACCATATACTTTAGAAGAACAGA

#### **SEQ ID NO: 712**

- 20 za24e08.s1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:293510 3', mRNA sequence gi|1225735|gb|N69574.1|N69574[1225735]
- W WAACTAATATTAAATAGTAAATTTAATGTGTATTAATATTGTCATATAATATTGNN A MAAMAATTACTCATGTAAATGTAAATATTACATTGAGGATATAGTAAATATTAAATTTAC W WAATGTCATTGAGGACAGTATTTCAAACTAGCTTTTTTAAAAAGAAAAACAGAAGA
- 25 TGGCAGTGAATAGAACAGTGATTGTTCATACTACTTGGATCTACTGCCTTAATTT ATACTAGGATGTCAATCCACCATTGATTTTGGACCATCAGTGCCAATGTCNACGT AGCCAAAAAGGCCAAT

#### **SEQ ID NO: 713**

- Human mRNA for gamma-interferon inducible early response gene (with homology to platelet proteins) gi|33917|emb|X02530.1|HSINFGER[33917]

  GAGACATTCCTCAATTGCTTAGACATATTCTGAGCCTACAGCAGAGGAACCTCCA
  GTCTCAGCACCATGAATCAAACTGCGATTCTGATTTGCTGCCTTATCTTTCTGACT
  CTAAGTGGCATTCAAGGAGTACCTCTCTCTAGAACCGTACGCTGTACCTGCATCA
- 35 GCATTAGTAATCAACCTGTTAATCCAAGGTCTTTAGAAAAACTTGAAATTATTCC
  TGCAAGCCAATTTTGTCCACGTGTTGAGATCATTGCTACAATGAAAAAGAAGGGT
  GAGAAGAGATGTCTGAATCCAGAATCGAAGGCCATCAAGAATTTACTGAAAGCA
  GTTAGCAAGGAAATGTCTAAAAGATCTCCTTAAAACCAGAGGGGAGCAAAATCG
  ATGCAGTGCTTCCAAGGATGGACCACACAGAGGCTGCCTCTCCCATCACTTCCCT
- 40 ACATGGAGTATATGTCAAGCCATAATTGTTCTTAGTTTGCAGTTACACTAAAAGG
  TGACCAATGATGGTCACCAAATCAGCTGCTACTACTCCTGTAGGAAGGTTAATGT
  TCATCATCCTAAGCTATTCAGTAATAACTCTACCCTGGCACTATAATGTAAGCTCT
  ACTGAGGTGCTATGTTCTTAGTGGATGTTCTGACCCTGCTTCAAATATTTCCCTCA
  CCTTTCCCATCTTCCAAGGGTACTAAGGAATCTTTCTGCTTTGGGGTTTATCAGAA
- 45 TTCTCAGAATCTCAAATAACTAAAAGGTATGCAATCAAATCTGCTTTTTAAAGAA TGCTCTTTACTTCATGGACTTCCACTGCCATCCTCCCAAGGGGCCCAAATTCTTTC AGTGGCTACCTACATACAATTCCAAACACATACAGGAAGGTAGAAATATCTGAA AATGTATGTGTAAGTATTCTTATTTAATGAAAGACTGTACAAAGTATAAGTCTTA GATGTATATATTTCCTATATTGTTTTCAGTGTACATGGAATAACATGTAATTAAGT

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**SEQ ID NO: 714** 

ab21g06.r1 Stratagene lung (#937210) Homo sapiens cDNA clone IMAGE:841498 5' similar to gb:X54304 MYOSIN REGULATORY LIGHT CHAIN 2, NONSARCOMERIC (HUMAN);, mRNA sequence gi|2217534|gb|AA487370.1|AA487370[2217534]

10 ACAAGGAAGATTTGCATGATATGCTTGCTTCTCTAGGGAAGAATCCCACTGATGC
 ATACCTTGATGCCATGATGAATGAGGCCCCAGGGCCATTCAATTTCACCATGTTC
 CTGACCATGTTTGGTGAGAAGTTAAATGGCACAGATCCTGAAGATGTCATCAGA
 AACGCCTTTGCTTTGATGAAGAAGCAACAGGCACCATTCAGGAAGATTACC
 TAAGAGAGCTGCTGACAACCATGGGGGATCGGTTTACAGATGAGGAAGTGGATG
 AGCTGTACAGAGAAGCACCTATTGACAAAAAAGGGGAATTTCAATTACATCGAGT
 TCACACGCATCCTGAAACATGGAGCCAAAGACAAAGATGACTGAAAGAACTTTA
 G

**SEQ ID NO: 715** 

20 H.sapiens mRNA for central cannabinoid receptor gi|736236|emb|X81120.1|HSCANN6[736236]

- TCTGGTGCTGTGCGTCATCCTCACTCCGCAGCCTCCGCTGCAGGCCTTCCTAC
  CACTTCATCGGCAGCCTGGCGGTGGCAGACCTCCTGGGGAGTGTCATTTTTGTCT
  ACAGCTTCATTGACTTCCACGTGTTCCACCGCAAAGATAGCCGCAACGTGTTTCT
  GTTCAAACTGGGTGGGGTCACGGCCTCCTTCACTGCCTCCGTGGGCAGCCTGTTC
- 35 CTCACAGCCATCGACAGGTACATATCCATTCACAGGCCCCTGGCCTATAAGAGGA TTGTCACCAGGCCCAAGGCCGTGGTGGCGTTTTGCCTGATGTGGACCATAGCCAT TGTGATCGCCGTGCTCCCTCCTGGGCTGGAACTGCGAGAAACTGCAATCTGTT TGCTCAGACATTTTCCCACACATTGATGAAACCTACCTGATGTTCTGGATCGGGG TCACCAGCGTACTGCTTCTGTTCATCGTGTATGCGTACATGTATATTCTCTGGAAG
- 45 GTGAACCCCATCATCTATGCTCTGAGGAGTAAGGACCTGCGACACGCTTTCCGGA GCATGTTTCCCTCTTGTGAAGGCACTGCGCAGCCTCTGGATAACAGCATGGGGGA CTCGGACTGCCTGCACAAACACGCAAACAATGCAGCCAGTGTTCACAGGGCCGC AGAAAGCTGCATCAAGAGCACAGTCAAGATTGCCAAGGTAACCATGTCTGTGTC CACAGACACGTCTGCCGAGGCTCTGTGAGCCTGATGCCTCCCTGGCAGCACAGG

**SEO ID NO: 716** 

Human mRNA for dihydropteridine reductase (hDHPR)

- 15 gi|30818|emb|X04882.1|HSDHPR[30818] CGGAGCCGGCTGCAGGAGCAGGATGCCGCCGCCGCCGCTGCAGGCGAGGC GCGCCGGGTGCTGTGCGCGCGCAGGGGCGCTCTGGGTTCTCGATGCGTGCA GGCTTTTCGGGCCCGCAACTGGTGGGTTGCCAGCGTTGATGTGGTGGAGAATGAA GAGGCCAGCGCTACGATCATTGTTAAAATGACAGACTCGTTCACTGAGCAGGCT 20 GACCAGGTGACTGCTGAGGTTGGAAAGCTCTTGGGTGAAGAAGGTGGATGCA ATTCTTTGCGTTGCTGGAGGATGGGCCGGGGGCAATGCCAAATCCAAGTCTCTCT \*TTAAGAACTGTGACCTGATGTGGAAGCAGAGCATATGGACATCGACCATCTCCA AAAGGCTGCCTGGATGGGACTCCTGGTATGATCGGGTACGGCATGGCCAAGGG 25 TGCTGTTCACCAGCTCTGCCAGAGCCTGGCTGGGAAGAACAGCGGCATGCCGCC CGGGGCAGCCGCCATCGCTGTGCTCCCGGTTACCCTGGATACCCCGATGAACAGG AAATCAATGCCTGAGGCTGACTTCAGCTCCTGGACACCCTTAGAATTCCTAGTTG AAACTTTCCATGACTGGATCACAGGGAAAAACCGACCGAGCTCAGGAAGCCTAA TCCAGGTGGTAACCACAGAAGGAAGGACGGAACTCACCCCAGCATATTTTTAGG 30 CCTCATCTCAGTGCCTATGAGGGGCCTGCCAGAAAAGTCACTAACCTGTCTCAGT GTGGCCTTGTCCAGCCTTGTGTTTTCTGTAACCCCTGTTTGTGGTACGAGATAATG AGTCCTATTTTCTCTCACATAATATGCATTTGCTCTCCTAGGACAGTGTAATACA TTTATGTGAAGTAAAGACATGCGAGACTGGTGGCCTGCAAATAGCATCCGTCAAT
- CTGTGTTAACTGCATAGGGAGGGCTCTGCATAGCACCTGCTATAGCGGTGTCATG
  35 TTGGATCGCTTTTGTGACTGTTCATCTGTCCTTGACAGTGGCTGTCATCTTGACTA
  CTTTGTTGATTTGTTGGTATTGGGGACATTTTAAAGGCTGAGTTATTTTTGAATGT
  CATGTTTATGTCATAGACGTAGTTTTCGCATCCTTGAATTAAACTGCCTTAACTCC
  TTTTGTGGTAT

AACATCTTGCAGACAGAGTGGCCATGTATGTGCATGCATATACACTCTACAGTGC TGTTAGACCTTTTGGGCTGCAGTTTCA

**SEQ ID NO: 718** 

5 zx10e07.s1 Soares\_total\_fetus\_Nb2HF8\_9w Homo sapiens cDNA clone IMAGE:786084 3', mRNA sequence gi|2162337|gb|AA448667.1|AA448667[2162337] ATAAATCTATAGTTTTATTAAGACAAAAACTGACAATGTAGTATGAAGTTTACAT TTAAA

**SEQ ID NO: 719** 

Human hyaluronate receptor (CD44) gene, exon 1 gi|180127|gb|M69215.1|HUMSCG01[180127]

- - 25 GAGCATGTGTGGAGAGAGGTGCCCATTCACACTGGCTTGAACACATGGGTTA GCTGAGCCAAATGCCAGCCCTATGACAGGCCATCAGTAGCTTTCCCTGAGCTGTT CTGCCAAGAAGCTAAAATTCATTCAAGCCATGTGGACTTGTTATTGAGGGGAAA AAGAATGAGCTCTCCCTCTTTCCACTTGGAAGATTCACCAACTCCCCACCCCTCA CTCCCCACTGTGGGCACGGAGGCACTGCGCCACCCAGGGCAAGACCTCGCCCTCT
  - 30 CTCCAGCTCCTCTCCCAGGATATCCAACATCCCTGTGAAACCAGAGATCTTGCTC CAGCCGGATTCAGAGAAATTTAGCGGGAAAGGAGAGGCCAAAGGCTGAACCCA ATGGTGCAAGGTTTACGGTTCGGTCATCCTCTGTCCTGACGCCGCGGGGCCAGC GGGAGAAGAAAGCCAGTGCGTCTCTGGGCGCAGGGGCCAGTGGGGCTCGGAGG CACAGGCACCCCGCGACACTCCAGGTTCCCCGACCCACGTCCCTGGCAGCCCCGA

  - 40 CCAGGGATCCTCCAGCTCCTTTCGCCCGCGCCCTCCGTTCGCTCCGGACACCATG GACAAGTTTTGGTGGCACGCAGCCTGGGGACTCTGCCTCGTGCCGCTGAGCCTGG CGCAGATCGGTGAGTGCCCGCCGCAGGCTGGGCAGCAAGATGGGTGCGGGGTGC TCAGCGCGGAC
  - 45 SEQ ID NO: 720

yi63g06.r1 Soares placenta Nb2HP Homo sapiens cDNA clone IMAGE:143962 5', mRNA sequence gi|851402|gb|R76770.1|R76770[851402] AATTCGGAACGAGGNCTGTACAACACAGTGTCATACAGGGATAATGCTATCATA TTTAATATGAAACAGTGTTACGGGCACAAATTACCCATTTCTACAAAATAAGTGT

GCAAGTGATGCCACATATTATCCATATTCAACTGAGCTGTCATCAAAATACATTT TATTTACAATATGTACTATGATCAGTTGGATATTAAGTTCTAAAATGATTTACTTC ACTGCTACATTATAAAGGTAAAAGCAATGTGTAGGAAAAAGTGTGAGATTGTGT TTTTACATACTGCTTTTGTAGTTGCCATCGCTGGTTCAGTTCGACTTATAACATAT GTCTTGCTTGTAGGATTTAACACCTCCAATAGGGGATTCTTCTAACATTACAGGA GGATTCTTAGGGGATCCGGGGCTTTTTCANCAGTATAT

#### **SEO ID NO: 721**

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15 TAGGATCCACAATAACAAGTTGATTCAGACTAATGTAGATATTTAGATTAGCAAG TATTGAACATTTGATTTCTTAGGACTGAGCTTTTAAATGAATTTCCATTATTTCTT CC

#### SEQ ID NO: 722

- Homo sapiens P2U nucleotide receptor mRNA, complete cds gi|984506|gb|U07225.1|HSU07225[984506]
- CGGCACGAGGCACCCCGAGAGGAGAAGCGCAGCGCAGTGGCGAGAGGAGCCCC

  TTGTGGCAGCAGCACTACCTGCCGAGAAAAATGCTGGAGGCTGGGCGTGGCCCC

  AGGCCTGGGACCTGTTTTTCCTGTTTCCCGCAGAGTTCCCTGCAGCCCGGTCCA

  CACCTGGAGAGCAGGGCTGGTCAGGGCGATGGCAGACCTGGGCCCCTGGA
  - 25 GGTCCAGGCGTGTGCATTCATGAGTGAGGAACCCGTGCAGGCGCTGAGCATCCT GACCTGGAGAGCAGGGCTGGTCAGGGCGATGGCAGCAGACCTGGGCCCCTGGA ATGACACCATCAATGGCACCTGGGATGGGGATGAGCTGGGCTACAGGTGCCGCT TCAACGAGGACTTCAAGTACGTGCTGCCTGTGTCCTACGGCGTGGTGTGCGT GCTTGGGCTGTGTCTGAACGCCGTGGCGCTCTACATCTTCTTGTGCCGCCTCAAG

  - 35 GCCGTGTGGGTGTTGGTGCTGGCCTGCCAGGCCCCCGTGCTCTACTTTGTCACCA CCAGCGCGCGGGGGCCGCGTAACCTGCCACGACACCTCGGCACCCGAGCTCT TCAGCCGCTTCGTGGCCTACAGCTCAGTCATGCTGGGCCTGCTCTTCGCGGTGCC CTTTGCCGTCATCCTTGTCTGTTACGTGCTCATGGCTCGGCGACTGCTAAAGCCAG CCTACGGGACCTCGGGCGCCTCCCTAGGGCCAAGTCCGTGCGCACCA
  - 40 TCGCCGTGGTGCTGTCTTCGCCCTCTGCTTCCTGCCATTCCACGTCACCCGC ACCCTCACTACTCCTTCCGCTCGCTGGACCTCAGCTGCCACACCCTCAACGCCAT CAACATGGCCTACAAGGTTACCCGGCCGCTGGCCAGTGCTAACAGTTGCCTTGAC CCCGTGCTCTACTTCCTGGCTGGGCAGAGGCTCGTACGCTTTGCCCGAGATGCCA AGCCACCCACTGGCCCAGCCCTGCCACCCCGGCTCGCCAGGCTGGGCCTGCG
  - 45 CAGATCCGACAGAACTGACATGCAGAGGATAGGAGATGTTTTGGGCAGCAGTGA GGACTTCAGGCGGACAGAGTCCACGCCGGCTGGTAGCGAGAACACTAAGGACAT TCGGCTGTAGGAGCAGAACACTTCAGCCTGTGCAGGTTTATATTGGGAAGCTGTA GAGGACCAGGACTTGTGCAGACGCCACAGTCTCCCCAGATATGGACCATCAGTG ACTCATGCTGGATGACCCCATGCTCCGTCATTTGACAGGGGCTCAGGATATTCAC

10 AA

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**SEQ ID NO: 723** 

aa50e04.s1 NCI\_CGAP\_GCB1 Homo sapiens cDNA clone IMAGE:824382 3', mRNA sequence

- 20 AACTTGAGCCAAGGGATAAATATAAGCAACCAATGGGCTGCAGGATAGTTGTAC
  AAAGTGTATCATGTATCTTCATAGCTTCTTTTGCCCATATAATGCATTCCACACTTA
  AGTTTCTCCTTCTAAAAGGGGACACGACAAGTTAATATGTCTCATAAATGTCTTA
  AATAAGTTGCATTTTCATGGCAAGCCCTCCACTGCCAGCAATGGATATACTCACAC
  CTATTGGAAAAAATCTAAAGTTAACAAACTGGTTTAGTATGGAAATGGTCTATTT
  - 25 GTTCCTCAGCTATGTTTCTGTATCCTACATTAGTGGCTCTCAGGAGG

**SEO ID NO: 724** 

HUMHBC4799 Human pancreatic islet Homo sapiens cDNA similar to alpha-1 antichymotrypsin, mRNA sequence gi|1262485|dbj|D83812.1|D83812[1262485]

- 35 CCCANAGACCCTGAAGCGGTGGAGAGACTCTCTGGAGTTCANAGAGATAGGTGA GCTCTACCTGCCAAAGTTTTCCANCTCGAGGGACTATAACCTGAACGACATNCTT CTCCAGCTGGGCATTGAGGAAGCCTTC

**SEQ ID NO: 725** 

zx84c12.s1 Soares ovary tumor NbHOT Homo sapiens cDNA clone IMAGE:810454 3', mRNA sequence gi|2179839|gb|AA457119.1|AA457119[2179839]
 CTCATCAAAACATGATTTATTAATTTTAAGCAAGAGTAAGCATATGTGATAGTGG CCAGCTTGGGGATAGAACTCTTCCTGGTTGATGCACAGTTCAGCACCTGTTGGGT CTTGGCTGTTGGGATAATTCTTTTTGGGTGAGGGGAACAGCCGTGGTCAAGGC
 TGCCTGCACCCCCATCCAGGCACAGGACCCTGGGCAAAGTCTCAAAAGAGGTAG TGTTTTTACTTTCGCACCAACAATACAACATAAGTATTGGGTACAAAAGAGGAGA TTTCCTTCCCCTCTACCTCAACGGGCAAAAGGCCTTCCATCTTCAGAAGAGGCTT GTGAGGACCATCGGTTGGATGACCTCCTAGTGAGTTCTGGCTCCCATTCAGAAGACA

CAGAGAAACCCACAAAAGGGGCCTGTGGATCTGGTTCCAGGTCTCAAGGGTACA GCTTGGTTACATCCCCAGGCCCC

**SEQ ID NO: 726** 

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- **SEO ID NO: 727**
- yr38g10.s1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:207618 3' similar to gb:L24038\_rna1 A-RAF PROTO-ONCOGENE SERINE/THREONINE-PROTEIN KINASE (HUMAN);, mRNA sequence
- 20 gi|1012590|gb|H59758.1|H59758[1012590]
- 25 CTCCCAAAATTTAGAAGTATCCCCAAAGCCAAGAGGAAACCAAATGATGGGAGG AGACAGGGGCTCAGTCTTTGGGCGGGGGTCCCCCAATTTCCAGAAGAACTGGG AAAAGGCACATGGGGGNCCCCCTTCATCTTCCCGGGGTGGGGGAATGGGGGGAT TCCTNAGGGCAGCNTCAGGGGCAGAGACGAACTTTGTTTGGGTTGGTNGGGCAA GGTTCCTTGGGCTTNGGAG

- **SEQ ID NO: 728**
- Human thyroid hormone receptor alpha 1 (TR-alpha-1) gene, complete cds gi|339662|gb|M24748.1|HUMTHRA1A[339662]
- 40 TCCATCCACCTATTCCTGCAAATATGACAGCTGCTGTGTCATTGACAAGATCAC CCGCAATCAGTGCCAGCTGTGCCGCTTCAAGAAGTGCATCGCCGTGGGCATGGCC ATGGACTTGGTTCTAGATGACTCGAAGCGGGTGGCCAAGCGTAAGCTGATTGAG CAGAACCGGGAGCGGCGGCGGAAGGAGGAGATGATCCGATCACTGCAGCAGCG ACCAGAGCCCACTCCTGAAGAGTGGGATCTGATCCACATTGCCACAGAGGCCCA

CTACGACCCTGAGAGCGACACCCTGACGCTGAGTGGGGAGATGGCTGTCAAGCG GGAGCAGCTCAAGAATGGCGGCCTGGGCGTAGTCTCCGACGCCATCTTTGAACT GGGCAAGTCACTCTCTGCCTTTAACCTGGATGACACGGAAGTGGCTCTGCTGCAG 5 AGAAGAGTCAGGAGCGTACCTGCTGGCGTTCGAGCACTACGTCAACCACCGCA CATGATCGGGGCCTGCCACGCCAGCCGCTTCCTCCACATGAAAGTCGAGTGCCCC ACCGAACTCTTCCCCCACTCTTCCTCGAGGTCTTTGAGGATCAGGAAGTCTAAA GCCTCAGGCGGCCAGAGGTGTGCGGAGCTGGTGGGGAGGAGCCTGGAGAGAA 10 CGTCCTTGGATAGATTCAGCTCCCACACACACCCCGCACTGCCCAGGTCCCTC CTCAGACCTCCAGCCCTGGGACAGGGCAAACAACTGAACTTGCTATGGAAAGGA CAGTGTGGGAGGCTGGGGAGCTGTGTCCTGCAGTTCCCAGGACCCCATCCTCTC AGAAGGTAGGGGAAGGCGGGGGGGATTGAGAAGGGACAAGCCACCTTGACCGT 15 AGGGGAAGGAGGAATGTGGGCTGGGGGAAGATGCCCTCAACTCACCCCCTCACA CACATGAGAGAGCCCCCACCCAGTTCCTTGGCCTAGGTCTCCCCTCCAGGCTG AGGGCCTCTCTACTTCCCCAGATGCCTGGGTGCAAAGAACGGCTTGGCTTGGCTC CTCCTCTGGAGGTTAAAATTTATAGTCATTCTAACTGCACTTGGAAACCAAGCAA GGGGAGAAGACAAATGAAGAAAAACT

20

#### **SEQ ID NO: 729**

de 40d05:s1 Gessler Wilms tumor Homo sapiens cDNA clone IMAGE:8982813'similar to gb:X53416 ENDOTHELIAL ACTIN-BINDING PROTEIN (HUMAN);, mRNA sequence gi gi 2432277 gb AA598978.1 AA598978[2432277]

TTTTTTTTAATGGAAGCAAAACTTTATTCCTCTTGGCTGGAGAAGAAGAACTAGT
GGGTGGTTGTGTACAGGACCCCCATCCCTCACCCCTCCCAGAACCAAAGAAGAC
AAGCAGCGCCACCAAATGGCTCCCTCTGCCCAAGTGAAAGCCGAGAGGTCAGCG
GCTGGCTGGGGAGGCAGGTGAGCGCACACGGCACAGGGCAGGGCGGCTGCAG
TGACAGGCGGGCCAGGGCCGGCCTGGGCCGGGTTGAGGGGAAGAGGGCGG
GGCTGCTTGGGTAGCGGGCAGGCTTGGGGGCTGCCGGCTGCCCCAG
ACTCAGGGCACCACAACGCGGTAGGGGCTGCCTGGGATGTGCTCGTCCCCCCATT
TGACCACCAGTGTGTAATCCCCCTTGTCCTTGAGCAGGTAGGACACGCTGTAGAG
CCGGATTGCCAAGTTCTTTACCAGGAGTTTCCCGCAGGGGCCTTTTGCCATTAA
CCCCACC

- **SEQ ID NO: 730**
- yr86d03.s1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:212165 3' similar to gb:Z22548 THIOL-SPECIFIC ANTIOXIDANT PROTEIN (HUMAN);, mRNA sequence gi|1030355|gb|H68845.1|H68845[1030355]
- 40 TTCCCTAATACTTTATTGGNTACCTCTAGGCCTGTGTGCGGCTGGGTGGGCTTGG
  GGGAGGGCGTCACTATTCAGCTTCTAGGTGGAGGCATGAGAAGGCCTTGGCTAG
  NCCCTCCAGGGTCCCATACTGTGGAGTTTGGAGGGCAGGTCTGGCCTTTCCTGG
  GTCAGCATAGGGCACCCAGGTNGGGGCACAGGTGGACACCCAGCACAGGCACCT
  AGGCAGGGCACAAGCTCACTATCCGTTAGCCAGCCTAATTGTGTTTGGAGAAAT
  45 ATTCCTTGCTGTCATCCACGTTGGGCTTAATCGTGTCACTACCAGGCTTCCAGCCA
- 45 ATTCCTTGCTGTCATCCACGTTGGGCTTAATCGTGTCACTACCAGGCTTCCAGCCA GCGGGANAAACTTTCCCCATGCTCGTCTGTGTACTGGGAAGGNCTGGGACCAGC CGCAGAGCCTANATTCCACGGAGCGTCCCACAGGCAAAT

SEQ ID NO: 731

ab23b05.r1 Stratagene lung (#937210) Homo sapiens cDNA clone IMAGE:841617 5' similar to TR:E183625 E183625 ORNITHINE DECARBOXYLASE ANTIZYME;, mRNA sequence

**SEQ ID NO: 732** 

Human elastase III B mRNA, complete cds, clone pCL1E3

- 15 gi|607029|gb|M18692.1|HUMELA3A[607029]
  CCTATCATCGCAAAACTCATGATGCTCCGGCTGCTCAGTTCCCTCCTTGTGGC
  CGTTGCCTCAGGCTATGGCCCACCTTCCTCTCGCCCTTCCAGCCGCGTTGTCAATG
  GTGAGGATGCGGTCCCCTACAGCTGGCCCTGGCAGGTTTCCCTGCAGTATGAGAA
  AAGCGGAAGCTTCTACCACACCTGTGGCGGTAGCCTCATCGCCCCCGACTGGGTT
- 20 GTGACTGCCGCCACTGCATCTCGAGCTCCCGGACCTACCAGGTGGTGTTGGGCG
  AGTACGACCGTGCTGTGAAGGAGGGCCCCGAGCAGGTGATCCCCATCAACTCTG
  AGGACCTCTTTGTGCATCCACTCTGGAACCGCTCGTGTGTGCCTGTGGCAATGA
  AGGATCGCCCTCATCAAGGTGTGACACCCCCAGCTGGGAGACGCCGTCCAGCTC

**SEQ ID NO: 733** 

SEQ ID NO: 734

**CCAAAG** 

45

yv19b06.s1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:243155 3', mRNA sequence gi|1102102|gb|H94469.1|H94469[1102102] GCAAAACATTTATTCTTTTAAAAAAATCTATATACATTGCCATACAAAGATAC

CACATTGAAGCAGTTCTCAGGAACCTTCCAGTGAGCCTTCTCTTATAATTGCCCG AGCAAGATTTCGTGCCAGAGAAAGTCTCAGCATTTCCACCTTGGTGTNCTCTATG TCATCATCCTGGAGCTGCTCGGTATCAGATTCTCCATGCACAGGTCTTCTTGACGT CAAGTCCTCCAGACACCGCATCAACTCATAAGTCTGTTCTGCTGAGAAAATCACC TGTTTCTGTTCCAAAAAGGGGCAAGGCATCTGTCAGCAGAGTTCATCCCAGAAAGA CCGAAGGGGCAATCCGAGACGTCATCAAGGACAGAAGGA

#### **SEQ ID NO: 735**

5

aa91g07.s1 Stratagene fetal retina 937202 Homo sapiens cDNA clone IMAGE:838716 3'
 similar to TR:G173234 G173234 RIBOSOMAL 5S RNA-BINDING PROTEIN ;, mRNA sequence

gi|2180364|gb|AA457644.1|AA457644[2180364]

- 15 CCAAACCAAATGGATATCTGCTTTTAAGATTAGAATTTGTTCTTCATCCTTAAAGC AGAACTCATTGAGATGAAAAGATGCTCTTAATTTATCACAGAACTGTGTATTTAA TAGTATGCTTATTAAAAATCACGAAGTGTACTGGAATGCTAAGATAAAAGAACTGT ATAGTTTCTGTTATGTAATACGAGAATAGAAATGTTATTAAAATCTTTCTATAATT TCCAGTGCTTCTGTTTTGAAGAACAAAGGCTTAATCCCCAAGAGGAAGTAGATAT
- 20 GCCAGTGTTTTCTACATTGATCCTGAATTTGCTGAAGATCCA

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CTEECEGCAGETCCTGCGCCGAGTGCETGAAAAAGGGCCCETTTGGGAA 25 GAACTGCAGCGCGCGTGTCCGGGCCTGCAGCTGTCGAACAACCCCGTGAAGGG CAGGACCTGCAAGGAGAGGGACTCAGAGGGCTGCTGGGTGGCCTACACGCTGGA GCAGCAGGACGGATGGACCGCTACCTCATCTATGTGGATGAGAGCCGAGGTGA GGCCGC

#### 30 SEQ ID NO: 737

- 40 ATATTCCCCTCAGGTTCCCGGTTTCCATTTTGTT

#### **SEQ ID NO: 738**

zx35f11.s1 Soares\_total\_fetus\_Nb2HF8\_9w Homo sapiens cDNA clone IMAGE:788493 3', mRNA sequence gi|2166225|gb|AA452556.1|AA452556[2166225]

TGAACACTGAAAAGAACAATATATATATACTGTAAAATATGATGAATAAACCAAATG TAGCTATAAGAATCTTAAAGGATGATTATAGAAAAGGGA

**SEQ ID NO: 739** 

15 SEO ID NO: 740

- ye40b03.r1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:120173 5', mRNA sequence gi|734317|gb|T95693.1|T95693[734317]
- - 25 ATGGANTTCAGGAGGGGGGCCTTAAGGCCNTTCAGGCAGG

**SEQ ID NO: 741** 

Human (clone HSY3RR) neuropeptide Y receptor (NPYR) mRNA, complete cds gi|189313|gb|L01639.1|HUMNYRECA[189313]

- CGCATCTGGAGAACCAGCGGTTACCATGGAGGGGATCAGTATATACACTTCAGA TAACTACACCGAGGAAATGGGCTCAGGGGACTATGACTCCATGAAGGAACCCTG TTTCCGTGAAGAAAATGCTAATTTCAATAAAATCTTCCTGCCCACCATCTACTCC ATCATCTTCTTAACTGGCATTGTGGGCAATGGATTGGTCATCCTGGTCATGGGTT ACCAGAAGAAACTGAGAAGCATGACGGACAAGTACAGGCTGCACCTGTCAGTGG
- 35 CCGACCTCCTCTTTGTCATCACGCTTCCCTTCTGGGCAGTTGATGCCGTGGCAAAC
  TGGTACTTTGGGAACTTCCTATGCAAGGCAGTCCATGTCATCTACACAGTCAACC
  TCTACAGCAGTGTCCTCATCCTGGCCTTCATCAGTCTGGACCGCTACCTGGCCATC
  GTCCACGCCACCAACAGTCAGAGGCCAAGGAAGCTGTTGGCTGAAAAGGTGGTC
  TATGTTGGCGTCTGGATCCCTGCCCTCCTGCTGACTATTCCCGACTTCATCTTTGC
- 45 AAATCATCAAGCAAGGGTGTGAGTTTGAGAACACTGTGCACAAGTGGATTTCCA
  TCACCGAGGCCCTAGCTTTCTTCCACTGTTGTCTGAACCCCATCCTCTATGCTTTC
  CTTGGAGCCAAATTTAAAACCTCTGCCCAGCACGCACTCACCTCTGTGAGCAGAG
  GGTCCAGCCTCAAGATCCTCTCCAAAGGAAAGCGAGGTGGACATTCATCTGTTTC
  CACTGAGTCTGAGTCTTCAAGTTTTCACTCCAGCTAACACAGATGTAAAAGACTT

TTTTTTTATACGATAAATAACTTTTTTTTAAGTTACACATTTTTCAGATATAAAAG ACTGACCAATATTGTACAGTTTTTATTGCTTGTTTGGATTTTTGTCTTGTGTTTCTTT AGTTTTTGTG

5 SEQ ID NO: 742
>AA504554
CACCACGGTGACCGTTTTCATCAGCAGCTCCCTCAACACCTTCCGCTCCGAGAA
GCGATACAGCCGCAGCCTCACCATCGCTGAGTTCAAGTGTAAACTGGAGTTGCTG
GTGGGCAGCCCTGCTTCCTGCATGGAACTGGAGCTGTATGGAGTTGACGACAA

10 GTTCTACAGCAAGCTG GATCAAGAGGATGCGCTCCTGGGCTCCTACCCTGTAGATGACGGCTG

SEQ ID NO: 743 >M11723

- - 25 ACCGCCTGTGCCACTGCCCGGTGGGCTACACCGGACCCTTCTGCGACGTGGACAC CAAGGCAAGCTGCTATGATGGCCGCGGGCTCAGCTACCGCGGCCTGGCCAGGAC CACGCTCTCGGGTGCGCCCTGTCAGCCGTGGGCCTCGGAGGCCACCTACCGGAAC GTGACTGCCGAGCAAGCGCGGAACTGGGGACTGGGCGCCACGCCTTCTGCCGG AACCCGGACAACGACATCCGCCCGTGGTGCTTCGTGCTGAACCGCGACCGGCTG
  - 30 AGCTGGGAGTACTGCGACCTGGCACAGTGCCAGACCCCAACCCAGGCGCGCCT CCGACCCCGGTGTCCCCTAGGCTTCATGTCCCACTCATGCCCGCGCAGCCGGCAC CGCCGAAGCCTCAGCCCACGACCCGGACCCCGTCTCAGTCCCAGACCCCGGGAG CCTTGCCGGCGAAGCGGGAGCAGCCGCCTTCCCTGACCAGGAACGGCCCACTGA GCTGCGGGCAGCGGCTCCGCAAGAGTCTGTCTTCGATGACCCGCGTCGTTGGCGG
  - 35 GCTGGTGGCGCTACGCGGGGCGCACCCCTACATCGCCGCGCTGTACTGGGGCCA CAGTTTCTGCGCCGGCAGCCTCATCGCCCCCTGCTGGGTGCTGACGGCCGCTCAC TGCCTGCAGGACCGGCCCGCACCCGAGGATCTGACGGTGGTGCTCGGCCAGGAA CGCCGTAACCACAGCTGTGAGCCGTGCCAGACGTTGGCCGTGCGCTCCTACCGCT TGCACGAGGCCTTCTCGCCCGTCAGCTACCAGCACGACCTGGCTCTGTTGCGCCT
  - 40 TCAGGAGGATGCGGACGCAGCTGCGCGCTCCTGTCGCCTTACGTTCAGCCGGTG
    TGCCTGCCAAGCGGCGCCGCGCGCGCCCCCGAGACCACGCTCTGCCAGGTGGCC
    GGCTGGGGCCACCAGTTCGAGGGGGCGGAGGAATATGCCAGCTTCCTGCAGGAG
    GCGCAGGTACCGTTCCTCCCTGGAGCGCTGCTCAGCCCCGGACGTGCACGGAT
    CCTCCATCCTCCCCGGCATGCTCTGCGCAGGGTTCCTCGAGGGCGCACCGATGC
  - 45 GTGCCAGGGTGATTCCGGAGGCCCGCTGGTGTGTGAGGACCAAGCTGCAGAGCG CCGGCTCACCCTGCAAGGCATCATCAGCTGGGGATCGGGCTGTGGTGACCGCAA CAAGCCAGGCGTCTACACCGATGTGGCCTACTACCTGGCCTGGATCCGGGAGCA CACCGTTTCCTGATTGCTCAGGGACTCATCTTTCCCTCCTTGGTGATTCCGCAGTG

AGAGAGTGGCTGGGGCATGGAAGGCAAGATTGTGTCCCATTCCCCAGTGCGGCCAGCTCCGCCCAGGATGGCGCAGGAACTCAATAAAGTGCTTTGAAAATGCTG

**SEQ ID NO: 744** 

5 >S60489

CTACTCCTAGATATTTGGCATGATCTTCAGTATGATCTTGTGCTGTGCTATCCGCA GGAACCGCGAGATGGTCTAGA

**SEQ ID NO: 745** 

10 >M59916

15 TGGCGCTGGCGCTGGCTCTGTCTGACTCTCGGGTTCTCTGGGCTCCGGC
AGAGGCTCACCCTCTTTCTCCCCAAGGCCATCCTGCCAGGTTACATCGCATAGTG
CCCCGGCTCCGAGATGTCTTTGGGTGGGGGAACCTCACCTGCCCAATCTGCAAAG
GTCTATTCACCGCCATCAACCTCGGGCTGAAGAAGGAACCCAATGTGGCTCGCGT
GGGCTCCGTGGCCATCAAGCTGTGCAATCTGCTGAAGATAGCACCACCTGCCGTG
20 TGCCAATCCATTGTCCACCTCTTTGAGGATGACATGGTGGAGGTGTGGAGACGCT

20 TGCCAATCCATTGTCCACCTCTTTGAGGATGACATGGTGGAGGTGTGGAGACGCT
CAGTGCTGAGCCCATCTGAGGCCTGTGGCCTGCTCCTGGGCTCCACCTGTGGGCA
CTGCGGACATTTCTCATCTTGGAACATCTCTTTGCCTACTGCCGAAGCCGCCCGAGCCCCAGGTGCCCCTGTCAGCCGCATCCTCTCCT
CACTGACCTGCACTGGGATCATGACTACCTGGAGGGCACCGGACCCTGACTGTGC

45 GCCGTCTGGCTACTCTTTGTGCCCAGCTCTCTGCCCGTGCTGACAGCCCTGCTCTG
TGCCGCCACCTGATGCCAGATGGGAGCCTCCCAGAGGCCCAGAGCCTGTGGCCA
AGGCCACTGTTTTGCTAGGGCCCCAGGGCCCACATTTGGGAAAGTTCTTGATGTA
GGAAAGGGTGAAAAAGCCCAAATGCTGCTGTGGTTCAACCAGGCAAGATCATCC
GGTGAAAGAACCAGTCCCTGGGCCCCAAGGATGCCGGGGAAACAGGACCTTCTC

**SEQ ID NO: 746** 

>W74362

TGAAGATGGAGCTAATCTTTCCTCTGCTCGTGGCATTTTGTCGCTTATCCAGTCTT

10 CTACTCGTAGGGCATACCAGCAGATCTTGGATGTGCTGGATGAAAATCGCAGAC
CTGTGTTGCGTGGTGGGTCTGCCGCCACTTCTAATCCTCATCATGACAACGT
NAGGTATGGCATTTCAAATATAGATACAACCATTGAAGGAAAGACCCCCNCNCC
NCGACTGTNNTAGATGCANCN

CCCCCCAGAAGACAGATAATCAAACTAAATAGACGTCTA

15

5

**SEQ ID NO: 747** 

>N71365

SEQ ID NO: 748 >AA454662

**SEQ ID NO: 749** 

>AA450180

SEQ ID NO: 750 >N76338

**SEQ ID NO: 751** 

15 >M60626 CCCAGAGCAAGACCACAGCTGGTGAACAGTCCAGGAGCAGACAAGATGGAGAC AAATTCCTCTCCCCACGAACATCTCTGGAGGGACACCTGCTGTATCTGCTGGC TATCTCTTCCTGGATATCATCACTTATCTGGTATTTGCAGTCACCTTTGTCCTCGG 20 AGTCACCACCATCAGTTACCTGAACCTGGCCGTGGCTGACTTCTGTTTCACCTCC ACTTTGCCATTCTTCATGGTCAGGAAGGCCATGGGAGGACATTGGCCTTTCGGCT GGTTCCTGTGCAAATTCCTCTTTACCATAGTGGACATCAACTTGTTCGGAAGTGTC THE CTGATCGCCCTCATTGCTCTGGACCGCTGTGTTTGCGTCCTGCATCCAGTCTG 25 《GGTGATGGCTCTGCTCACATTGCCAGTTATCATTCGTGTGACTACAGTACCTG GTAAAACGGGGACAGTAGCCTGCACTTTTAACTTTTCGCCCTGGACCAACGACCC TAAAGAGAGGATAAATGTGGCCGTTGCCATGTTGACGGTGAGAGGCATCATCCG GTTCATCATTGGCTTCAGCGCACCCATGTCCATCGTTGCTGTCAGTTATGGGCTTA TTGCCACCAAGATCCACAAGCAAGCTTGATTAAGTCCAGTCGTCCCTTACGGGT 30 CCTCTCCTTTGTCGCAGCAGCCTTTTTTCTCTGCTGGTCCCCATATCAGGTGGTGG CCCTTATAGCCACAGTCAGAATCCGTGAGTTATTGCAAGGCATGTACAAAGAAAT TGGTATTGCAGTGGATGTGACAAGTGCCCTGGCCTTCTTCAACAGCTGCCTCAAC  ${\tt CCCATGCTCTATGTCTTCATGGGCCAGGACTTCCGGGAGAGGCTGATCCACGCCC}$ TTCCCGCCAGTCTGGAGAGGGCCCTGACCGAGGACTCAACCCAAACCAGTGACA 35 CAGCTACCAATTCTACTTTACCTTCTGCAGAGGTGGCGTTACAGGCAAAGTGAGG AGGGAGCTGGGGGACACTTTCGAGCTCCCAGCTCCAGCTTCGTCTCACCTTGAGT TAGGCTGAGCACAGGCATTTCCTGCTTATTTTAGGATTACCCACTCATCAGAAAA AAAAAAAAGCCTTTGTGTCCCCTGATTTGGGGAGAATAAACAGATATGAGTTT ATTATTGACTTCTTTTTTGATTTTGGACCTCAGCCTCGGGTGGTCAGGGTGGGAAA 40 TGATAGGAAGAAGCTGTCATCTGCATCCTAGTTTGCCTGAAATGAACCCAAATAA TACCCATTATTATTAGTCCTGAATTATGAGTAGTGAATGATACCCATCATTCTGGC ATCATGATGAGTAGTGTCCACTTCCATTCTGAAAAGTGCCCTGCTGTGAAAAATA AATTATATAGTCATCCTAGGTAAATGAAGGAGGAGGGAGAAGTGTGAAAGAGTA TGGCTTAAATCAGACAAGATATACAAGAAGATACTTTATATAGGGCAGGAGCGG 45 GAGGTCAGGAATTCGAGAACAGCCTGGCCAACATGGTGAAACCCTGTCTCTACT AAAAATACAAAAATTAGCTGGGCGTAGTGGCAGGCTCCCGTAATCCCAGCTACT CAGGAGACCGAGGCAGGAGATCGCTTGGACCTGGAAGGCGGAGGTTGTAGTGA

SEQ ID NO: 752 >X70070

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45

TCAAGCTCGCCCGCGCAGCCGAGCCGGGCTGGGCGCTGTCCTCGGGGGCCTG GGGAACCGCGCGTTTGGAGATCGGAGGCACCTGGAACCCGTGGCAAGCGCCGA GCCGGGAGACAGCCGAGGAACCACGGGTTCTGGAGCTAGGAGCCGGAAGCTG GGAGTCCGGAGGAGCCGGAGCCCGGAGCCCGGGGCGCGCGTCTG 10 GGTCTGGCGCTTCCCGACTGGACGGCGCCCCCCTGGTCTTCGCCACGCGCCCTC CCCTGGGCTCGCGTTCATCGGTCCCGCCTGAGACGCGCCCACTCCTGCCCGGAC TTCCAGCCCGGAGGCGCCGGACAGAGCCGCGGACTCCAGCGCCCACCATGCGC CGGGCGCAGGCCGGACTGGAGGAGGCGCTGCTGGCCCCGGGCTTCGGCAACGCT 15 TCGGGCAACGCGTCGGAGCGCGTCCTGGCGGCACCCAGCAGCGAGCTGGACGTG AACACCGACATCTACTCCAAAGTGCTGGTGACCGCCGTGTACCTGGCGCTCTTCG TGGTGGCACGGTGGCAACACGGTGACGCGTTCACGCTGGCGCGGAAGAAGT CGCTGCAGAGCCTGCAGAGCACGGTGCATTACCACCTGGGCAGCCTGGCGCTGT CCGACCTGCTCACCCTGCTGGCCATGCCCGTGGAGCTGTACAACTTCATCTG 20 GGTGCACCACCCCTGGGCCTTCGGCGACGCCGGCTGCCGCGGCTACTACTTCCTG CGCGACGCCTGCACCTACGCCACGGCCTCAACGTGGCCAGCCTGAGTGTGGAG MGCCCTACCTGGCCATCFGCCACCCCTTCAAGGCCAAGACCCTCATGTCCCGAAGCC TATATGCTGTTCACCATGGGCGAGCAGAACCGCAGCGCCGACGCCAGCACGCCGG ○ CGGCCTGGTGTGCACCCCCACCATCCACACTGCCACCGTCAAGGTCGTCATACAG 25 GTCAACACCTTCATGTCCTTCATATTCCCCATGGTGGTCATCTCGGTCCTGAACAC CATCATCGCCAACAAGCTGACCGTCATGGTACGCCAGGCGGCCGAGCAGGGCCA AGTGTGCACGGTCGGGGGCGAGCACAGCACATTCAGCATGGCCATCGAGCCTGG CAGGGTCCAGGCCTGCGGCACGGCGTGCGCGTCCTACGTGCAGTGGTCATCGCC 30 TTTGTGGTCTGCCTGCCCTACCACGTGCGGCGCCTCATGTTCTGCTACATCTC GGATGAGCAGTGGACTCCGTTCCTCTATGACTTCTACCACTACTTCTACATGGTG ACCAACGCACTCTTCTACGTCAGCTCCACCATCAACCCCATCCTGTACAACCTCG TGGCGCGCAGGAGGAAGAGCCAGCCTTCTCGAGGAAGGCCGACAGCGTGTCC 35 AGCAACCACCCTCTCCAGCAATGCCACCCGCGAGACGCTGTACTAGGCTGTGC GCCCGGAACGTGTCCAGGAGGAGCCTGGCCATGGGTCCTTGCCCCCGACAGAC AGAGCAGCCCCACCCGGGAGCCTTGATGGGGGTCAGGCAGAGGCCAGCCTGCA CTGGAGTCTGAGGCCTGGGACCCCCCCCCCCCCCCCCCACCCCCTAACCCATGTTTCTCATT AGTGTCTCCCGGGCCTGTCCCCAACTCCTCCCCACCCCTCCCCATCTCTTTG 40 AAAGCCAGAACAAGAGCGCTCCTCTCCCAGATAGGAAAAGGGCCTCTAACAA GGAGAAATTAGTGTGCGGCAAAAGGCAGTTTTCTTTGTTCTCAGACTAATGGATG GTTCCAGAGAAGGAAATGAAATGTGCTGGGTGGGCCGGGCCTCCGGCGGCCCG

GCTGCTGTTCCCATGTCCACATCTCTGAGGCCTGCACCCCCTCTGTCTAGCTCGGG GAGTCCAGCCCCAGTCCCGCAGGCTCCGTGGCTTTGGGCCTCACGTGCAGACCCT

GCTCAGGCCTCAAGATCTTCAGCTGTGGCCTCTCGGGCTCGGCAGAAGG GACGCCGGATCAGGGCCTGGTCTCCAGCACCTGCCCGAGTGGCCGTGGCCAGG ATGGGGTGCGCATTCCGTGTGCTTTGCTTGTAGCTGTGCAGGCTGAGGTCTGGAG CCAGGCCCAGAGCTGGCTTCAGGGTGGGCCTTGAGAAGGGGAATGTGGGACAG 5 GGGCGATGGTGCCTGGTCTCTGAGTAAGATGCCAGGTCCCAGGAACTCAGGCTTC AGGTGAGAAGGAGCGGTGTGTCCAGGCACCGCTGGCCGGCAGCCCTGGGCTGAG GCACAGACTCATTTGTCACCTTCTGGCGGCGGCAGCCCTGGCCCCGGCCTCCAAG CAGTTGAAAAAGCTGGCGCCTCCTTGGTCTCTAGGATCCAGGCTCCACAGAGCAC ATGACTAGCCAGGCCCCTGGCTTAAGAAGGTCGCCTAAGCCTAAGAGAAGACAG 10 TCCCAGGAGAAGCTGGCCGGGACCAGCCAGGAGCTGGGAGCCACAGGAAGCAA AAGTCAGCCTTTCCTCAAGGGATTTCCCTGTCTCAGAGCAGCCTTTGCCCCAGG GAAATGGGCTCTGGGCTGCCTGCACCGGCCATGTCGACCCAGGACCCGGA CACCTGGTCTTGGGCTGTTCAGCCACTTTGCCTTCTCTGGACTCAGTTTCCCCG TCTGAGAAATGAGAGTCGAATGCTACAGTATCTGCAGTCGCTTGGATCTGGCTGT 15 TGAGTTGACGGGTTCCTTGAACCCCACAAAATCCCTCTCCAACCACAGGACCCTT CGGCTCACCAAGAACGGGGCCCAGGGGAGTCAGGCCTATTCGCTGCACTTCCTG CCAAACTTTGCCCCCACAAGCCTGGTCATCAGCCAGGCAGCCCTCCCAGTGCCCA AGGGCCACCAACCCCAGGGAAACAGGGCCAGCACAGAGGGGCCTTCCTCCCCA 20 GATGTCCAGAGGTCGGTGCAGCCCCTATCCCTGCTCAGGAGTGGGCTCAGAGTCT AGCAAATGCTAAGGCCCCTCAGGCTGGGCTCTGAACGAGGACCTGGACTCAGAG I COMPANY TOTAL CONTROL OF THE CONTR TCAGGATGGTGCTCTGAGAGAGGGCAGAGTGGATGCCCCACTGCCCTAGACCCT CGGTAGACGTGGGGTCTCTGGGGCGGGTCTGTGGCTGTGACTGAAGTCGGCTTT TCCATGCACCACAGACACCCACGACACCTGATCTCGTATCACTAGCTTGCGGC CAGGTCATGATGTGGCCCCGGAAGCTGGCCCTGCGTGCCATGAGTGCGTCGGTCA 30 TGGAGTCCGGAGCCCTGAGCCGGCCCTTGGTGACGGCACAGCCCTCACAGCTC CTCTCAATAAAGGTGGCCGAAGGGCCTCGATGTGG

**SEQ ID NO: 753** 

35 >X58454 ATGCTGCCGCCAGGCAGCAACGGCACCGCGTACCCGGGGCAGTTCGCTCTATAC CAGCAGCTGGCGCAGGGGAACGCCGTGGGGGGGCTCGGCGGGGGCACCGCCACTG GGGCCTCACAGGTGGTCACCGCCTGCTGACCCTACTCATCATCTGGACCC TGCTGGGCAACGTGCTGGTGCGCAGCCATCGTGCGGAGCCGCCACCTGCGCG 40 CCAACATGACCAACGTCTTCATCGTGTCTCTGGCCGTGTCAGACCTTTTCGTGGC GCTGCTGGTCATGCCCTGGAAGGCAGTCGCCGAGGTGGCCGGTTACTGGCCCTTT GGAGCGTTCTGCGACGTCTGGGTGGCCTTCGACATCATGTGCTCCACTGCCTCCA TCCTGAACCTGTGCGTCATCAGCGTGGACCGCTACTGGGCCATCTCCAGGCCCTT CCGCTACAAGCGCAAGATGACTCAGCGCATGGCCTTGGTCATGGTCGGCCTGGC 45 ATGGACCTTGTCCATCTCATCTCCTTCATTCCGGTCCAGCTCAACTGGCACAGG GACCAGGCGGCCTCTTGGGGCGGGCTGGACCTGCCAAACAACCTGGCCAACTGG ACGCCCTGGGAGGAGCTTTTGGGAGCCCGACGTGAATGCAGAGAACTGTGAC TCCAGCCTGAATCGAACCTACGCCATCTCTTCCTCGCTCATCAGCTTCTACATCCC CGTTGCCATCATGATCGTGACCTACACGCGCATCTACCGCATCGCCCAGGTGCAG

ATCCGCAGGATTTCCTCCCTGGAGAGGGCCGCAGAGCACGCGCAGAGCTGCCGG
AGCAGCGCGCGCCCGACACCAGCCTGCGCGCTTCCATCAAGAAGGAG
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5 CCTCCGGCCGGCTTCCCCTGCGTCAGTGAGACCACCTTCGACGTCTTCGTCTGGTT
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GGAGACGGTGAACATCAGCAATGAGCTCATCTCCTACAACCAAGACATCGTCTTC
CACAAGGAAATCGCAGCTGCCTACATCCACATGATGCCCAACGCCGTTACCCCCG
10 GCAACCGGGAGGTGGACAACGACGAGGAGGAGGTCCTTTCGATCGCATGTTCC
AGATCTATCAGACGTCCCCAGATGGTGACCCTGTTGCTGAGTCTTCTTCGGGAGCT
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ATTCCATTAA

15 SEQ ID NO: 754

>D13538

- TGGCTGCCGTGGTGGGCTTCCTCATCGTCTTCACCGTGGTGGGCAACGTGCTGGT GGTGATCGCCGTGCTGACCAGCCGGGCGCTGCGCGCGCCACAGAACCTCTTCCTG GTGTCGCTGGCCTCGGCCGACATCCTGGTGGCCACGCTGGTCATGCCCTTCTCGT ATGGCCAACGAGCTCATGGCCTACTGGTACTTCGGGCCATGTGTGCCGATGTGCCATCA
  - 25 GCCTGGACCGCTACTGGTCGGTGACGCAGGCCGTCGAGTACAACCTGAAGCGCA CACCACGCCGCGTCAAGGCCACCATCGTGGCCGTGTGGCTCATCTCGGCCGTCAT CTCCTTCCCGCCGCTGGTCTCGCTCTACCGCCAGCCCGACGGCGCCGCCTACCCG CAGTGCGGCCTCAACGACGAGACCTGGTACATCCTGTCCTCCTGCATCGGCTCCT TCTTCGCGCCCTCCTCATCATGGGCCTGGTCTACGCGCGCATCTACCGAGTGGC

  - 40 CTTCTGGATCGCTACTGCAACAGCTCGCTCAACCCGGTCATCTACACGGTCTTC AACCAGGATTTCCGGCGATCCTTTAAGCACATCCTCTTCCGACGGAGGAGAAGG GGCTTCAGGCAGTGACTC

**SEO ID NO: 755** 

45 >N76944

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**SEQ ID NO: 756** 

5 >AA451716

10 GCAGCTGTGATCCAGCAGCAGCTGGCAAAGCTTAGTAAGCAACCTCATCCCCAG ATGCATCCGCTCAGCCAGTGTTGTGATTGCTAGATACTATCTGTAAGTGAACCAA ACTAAAATTCATTTATGAACCAAGAAAGGAAGCCAAGTTGAAAAGGTCTCGAGT TAAATCGAGAATGATTCAGGCGGGCCGGCTCTCTGAGCA CCTTTGGATGCACTTCAGCTTCTGTCTTG

15

**SEQ ID NO: 757** 

>H19264

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- 20 GGTGGTGAACCTGGTNGTGGAGAGCACACCTACTTTAGCCAACTTGGGCAGGGT GGCCCAGGTCCTGAGGCTGATGCGGATCTTCCGATCTTAAAGCTGGCCAGGCACT

20 20 20 20 20 20 20

25 SEQ ID NO: 758

>AA598527

- 30 CCACCATTTGGCTAATATTATTTCATTAAAGACTGAATTTAGATTTTAGGAAATA
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  TGTTAAATTAAGAAGAAATTCATAGACACCATTTTTTTCCTGTTACAACATATGG
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- 35 TAAAGTTACTTTCTGTCCAGGTCGAAACATTGTTC

**SEQ ID NO: 759** 

>AA286908

SEQ ID NO: 760 >AA280924

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10 TGGTCAACACGCGGGCATCGCTTCAAGGTTGCTGATCCCACACCCTTTCATATT CAAGCTGAAGTGACGATGAAAACAAATTTCTTTGGTACCCGAGATGTGTGCACA GAATTACTCCCTCTAATAAAACCCCAAGGGAGAGTGGTGAACGTATCTAG

**SEQ ID NO: 761** 

15 >AA279601

AAAGATGTAAATCGTGGAGGTTACGGCCAAGCCAGTTCCATTAAATTCAGGAGT CAGATTCCAGGTTTATGTAGAAGTTTCTAAAATGAAAATCAATGTTACTGAAATT CCTGACACATTGCGTGAAGATCAAATGAGAGACAAACTAGAGCTGAGCTTTTCA 

20 GGGAGTGCAGTCATCACGGTTGT G

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>N22980

1.1 GTTAAAACATGAAAAAAATTTTATTGTTTTAGACAAAGAGGCCACTTTTGGAAA 25 ATAATACTTTTTTTTTTTTGAATCAGGTGAAGACAGAGTTAAAATCACATA GGATTTGCATTTTAAAAAAGGAAAGCACTAGGATTGTTGGCACTGGAGTAACTA TTTACACTGAACAGAGGTTTGGCCTTTTACATAACATCGATACAATGCATTTTCC AAAGTCTGAGAAATAACAAGGTTCTGTCTCGAATGCTTCACAGAGGAGGTTCGG ATTTGGGGACAAGTGTCATTAATGAGGGCCATGGAAGTTCGTCAGCTTCAGAGTC 30 ACATGCAATCTGATCCTGGGCGGTTCCCCNGCTGGGGAGCACTTGGCTACGGAAT

TGAAAGCTAATGGGGAGGGGTGGGGC

**SEQ ID NO: 763** >T61575

- 35 GATTATATCATGGTATATGAAGCACTGGTGAGGTCTATGTCACCAGAAATTCCCA GTTTGCTGATTTCATTGAGTTTTTTAACCCGATGATNGTACTGCAACAAGTNAGC ATNNGTCACTGCAACCNAACNNGNGGGGGGGGAAGGTNCACCCNNNNTTNTTTT TGAAAGGGTTCCCATTTTCNAANGGGGAAACCGNTNTTTTTCTTCCCTNCCCNGT TATTATCCAGCTTTGTATTGCAAACAATGACTCTCCTGTTGTTCTCATTGAAGCGT
- 40 GGGGTTAAAGTGGGAGGCAACATCATTCCCTCTTTGGGAAATCTAAGGCAATTC TGTTTGCATTGGGGCTTCACCGTGCCCAGAATTGTTATCAGCATGCGAGGGGACC ACTCCCCGGGGGAAAGGGCAGGGTTATAGGGGACAATCAGTGGGCCCGNAGG GGGNCCATGGGGNCCAGTGGCAGGGGNAGGGTNCCGTGGCNCTTGGCTTT

45 **SEQ ID NO: 764** 

>R23586

AGAACGAGCTGGACCAGAAGAAAGTAAAATATCCCAAAATGACAGACCTCAGC AAGGGTGTGATTGAGGAGCCCAAGTAGCGCCTGCTTNGCGTGGGTGGATCCAAC ACCAGCCTGCGTTCGTGGGACTTGCCTCAGATCAGCCTGCGACTGCAAGATTCT

SEQ ID NO: 765 >L08044

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44.

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**SEQ ID NO: 766** 

SEQ ID NO: 767 > U39613

ATGGCAGATGATCAGGGCTGTATTGAAGAGCAGGGGGTTGAGGATTCAGCAAAT 35 GAAGATTCAGTGGATGCTAAGCCAGACCGGTCCTCGTTTGTACCGTCCCTCTTCA GTAAGAAGAAGAAAATGTCACCATGCGATCCATCAAGACCACCCGGGACCGAG TGCCTACATATCAGTACAACATGAATTTTGAAAAGCTGGGCAAATGCATCATAAT AAACAACAAGAACTTTGATAAAGTGACAGGTATGGGCGTTCGAAACGGAACAGA CAAAGATGCCGAGGCGCTCTTCAAGTGCTTCCGAAGCCTGGGTTTTGACGTGATT 40 GTCTATAATGACTGCTCTTGTGCCAAGATGCAAGATCTGCTTAAAAAAAGCTTCTG AAGAGGACCATACAAATGCCGCCTGCTTCGCCTGCATCCTCTAAGCCATGGAGA AGAAAATGTAATTTATGGGAAAGATGGTGTCACACCAATAAAGGATTTGACAGC CCACTTTAGGGGGGATAGATGCAAAACCCTTTTAGAGAAACCCAAACTCTTCTTC ATTCAGGCTTGCCGAGGGACCGAGCTTGATGCCATCCAGGCCGACTCGGGG 45 CCCATCAATGACACAGATGCTAATCCTCGATACAAGATCCCAGTGGAAGCTGACT TCCTCTTCGCCTATTCCACGGTTCCAGGCTATTACTCGTGGAGGAGCCCAGGAAG AGGCTCCTGGTTTGTGCAAGCCCTCTGCTCCATCCTGGAGGAGCACGGAAAAGAC CTGGAAATCATGCAGATCCTCACCAGGGTGAATGACAGAGTTGCCAGGCACTTT

GAGTCTCAGTCTGATGACCCACACTTCCATGAGAAGAAGCAGATCCCCTGTGTGG
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**SEQ ID NO: 768** 

5 >H91337

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10 GGTCTAGTCCTTATACCGACTCAGATTCCTTAAGCATGCAGAGTCACTCGAATG AAAAAA

**SEO ID NO: 769** 

>M29870

20 TGTCCGTGCAAAGTGGTATCCTGAGGTGCGGCACCACTGTCCCAACACTCCCATC ATCCTAGTGGGAACTAAACTTGATCTTAGGGATGATAAAGACACGATCGAGAAA CTGAAGGAGAAGAAGCTGACTCCCATCACCTATCCGCAGGGTCTAGCCATGGCT

25 AGAAGAGAAGAAAATGCCTGCTGTTAA

SEQ ID NO: 770

>AA454652

TAAGTGGCAAATGCTATTCTAAGTGGCAAAACAATC

35

SEO ID NO: 771

>AA424315

AGGTCCCATTACAAAGCATGGTGAAATAGATTATGAAGCAATTGTGAAGCTTTCG GATGGCTTTAATGGAGCAGATCTGAGAAATGTTTGTACTGAAGCAGGTATGTTCG

40 CAATTCGTGCTGATCATGATTTTGTAGTACAGGAAGACTTCATGAAAGCAGTCAG
AAAAGTGGCTGATTCTAAGAAGCTGGAGTCTAAATTGGACTACAAACCTGTGTA
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AAAGTTAAAGAAAATAATGTATTGTATTGGCAATGATGTCATTAAAAGTATATGA
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45 AGAAATTGTATGTTTGTTAAAGTTGCATTTATTGCAGCAAG

**SEQ ID NO: 772** 

>AA460727

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TACATACTTGGTTCTCCAAATAACAACTGTGTGACATATAGCAGAAGGAATTAAG
GAATGCTGCACTTGTGATCCATACAAAACACCCAACATTTTAGGTTGTACATAATT

5 AGAGAAATATCTGAAACACTTTTTAAAACACTGTAGTAGCCAATACATAGAGGC
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CATAGGAACTACTCAATTTCTTTAAAAATCACTGAGCAAGAACAGCAACATTGAAC
TTTCATACTGATTTTACACAACTTCTATACAGTACCTTGACTTAAATCCAAGAGCA
AAAGTTAAGACTCTCCTCCTCTATTTTTGGTAAACCACTGCATGGTAAACTTAGA

10 TGACTCTTCCCCCTGGATTTTACCT

SEQ ID NO: 773 >L15189

CCTGCCTCGTACTCCTCCATTTATCCGCCATGATAAGTGCCAGCCGAGCTGCAGC 15 GGATAGCTGGAATGGCCTTAGTCATGAGGCTTTTAGACTTGTTTCAAGGCGGGAT TATGCATCAGAAGCAATCAAGGGAGCAGTTGTTGGTATTGATTTGGGTACTACCA ACTCCTGCGTGGCAGTTATGGAAGGTAAACAAGCAAAGGTGCTGGAGAATGCCG AAGGTGCCAGAACCACCCCTTCAGTTGTGGCCTTTACAGCAGATGGTGAGCGACT 20 TGTTGGAATGCCGGCCAAGCGACAGGCTGTCACCAACCCAAACAATACATTTTAT GCTACCAAGCGTCTCATTGGCCGGCGATATGATGATCCTGAAGTACAGAAAGAC A STANGARD AND A STAN SA GATGAAAGAGACTGCAGAAAATTACTTGGGGCACACAGGAAAAAATGCTGTGAT 125 CACAGTCCCAGCTTATTTCAATGACTCGCAGAGACAGGCCACTAAAGATGCTGGC CAGATATCTGGACTGAATGTGCTTCGGGTGATTAATGAGCCCACAGCTGCTGCTC TTGCCTATGGTCTAGACAAATCAGAAGACAAAGTCATTGCTGTATATGATTTAGG TGGTGGAACTTTTGATATTTCTATCCTGGAAATTCAGAAAGGAGTATTTGAGGTG AAATCCACAAATGGGGATACCTTCTTAGGTGGGGAAGACTTTGACCAGGCCTTGC 30 TACGGCACATTGTGAAGGAGTTCAAGAGAGAGACAGGGGTTGATTTGACTAAAG ACAACATGGCACTTCAGAGGGTACGGGAAGCTGCTGAAAAGGCTAAATGTGAAC TCTCCTCATCTGTGCAGACTGACATCAATTTGCCCTATCTTACAATGGATTCTTCT GGACCCAAGCATTTGAATATGAAGTTGACCCGTGCTCAATTTGAAGGGATTGTCA CTGATCTAATCAGAAGGACTATCGCTCCATGCCAAAAAGCTATGCAAGATGCAG 35 TGCCCAAGGTTCAGCAGACTGTACAGGATCTTTTTGGCAGAGCCCCAAGTAAAGC TGTCAATCCTGATGAGGCTGTGGCCATTGGAGCTGCCATTCAGGGAGGTGTGTTG GCCGGCGATGTCACGGATGTGCTCCTTGATGTCACTCCCCTGTCTCTGGGTAT TGAAACTCTAGGAGGTGTCTTTACCAAACTTATTAATAGGAATACCACTATTCCA 40 ACCAAGAAGAGCCAGGTATTCTCTACTGCCGCTGATGGTCAAACGCAAGTGGAA ATTAAAGTGTCAGGGTGAAAGAGAGATGGCTGGAGACAACAACTCCTTGGA CAGTTTACTTTGATTGGAATTCCACCAGCCCCTCGTGGAGTTCCTCAGATTGAAG TTACATTTGACATTGATGCCAATGGGATAGTACATGTTTCTGCTAAAGATAAAGG CACAAGACGTGAGCAGCAGATTGTAATCCAGTCTTCTGGTGGATTAAGCAAAGA 45 TGATATTGAAAATATGGTTAAAAATGCAGAGAAATATGCTGAAGAAGACCGGCG AAAGAAGGAACGAGTTGAAGCAGTTAATATGGCTGAAGGAATCATTCACGACAC AGAAACCAAGATGGAAGAATTCAAGGACCAATTACCTGCTGATGAGTGCAACAA GCTGAAAGAAGAGATTTCCAAAATGAGGGAGCTCCTGGCTAGAAAAGACAGTGA

AACAGGAGAAAATATTAGACAGGCAGCATCCTCTCTCAGCAGGCATCATTGAA

GCTGTTCGAAATGGCATACAAAAAGATGGCATCTGAGCGAGAAGGCTCTGGAAG TTCTGGCACTGGGGAACAAAAGGAAGATCAAAAGGAGGAAAAACAGTAATAAT AGCAGAAATTTTGAAGCCAGAAGGACAACATATGAAGCTTAGGAGTGAAGAGAC TTCC

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**SEQ ID NO: 774** 

>W60890

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- 10 CGGTGCTCGAATGATAACTATTGATGGGAAACAGATAAAACTTCAGATATGGGA TACGGCAGGGCAAGAATCCTTTCGTTCCATCACAAGGTCGTATTACAGAGGTGCA GCAGGAGCTTTACTAGTTTACGATATTACACGGAGAGATACATTCAACCACTTGA CAACCTGGTTAGAAGATGCCCGCCAGCATTCCAACTCCAACATGGTCATTATGCT TATTGGAAATAAAAGTGATTTAGAATCTAGAAGAAGAAGAAGAAGAAG
- 15 GTGAAGCTTTTGCACGAGAACATGGACTCATCTTCATGGAAACGTCTGCTAAGAC
  TGCTTCCAATGTAGAAGAGGCATTTATTAATACAGCAAAAGAAATTTATGAAAA
  AATTCAAGAAGGAGCTTTGACATTAATAATGAGGCCAATGGCATTAAAATTGGC
  CCTCAGCATNTGTTACCATGCCACACATGCAGGCNATCAGGGAGGCANCAGCTG
  GGGCNGCTCTGTTGANTCTGTTTATGCTANTGCCACGGGCTTCTCCCTTATCTTAN
  20 CCTTCCTCTGGNACTGGNTGACCTTTGAAAGGTTTGCCAGAGATTANCCGCAATC

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m T}$ 

SEQ ID NO: 775

>AA287196

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- 25 GTACACTGGTGTTGGACAGAGCAGCTTGGCTTTTCATGTGCCCACCTACTTACCT
  ACTACCTGCGACTTTCTTTTTCCTTGTTCTAGCTGACTCTTCATGCCCCTAAGATTT
  TAAGTACGATGGTGAACGTTCTAATTTCAGAACCAATTGCGAGTCATGTAGTGTG
  GTAGAATTAAAGGAGGACACGAGCCTGCTTCTGTTACCTCCAAGTGGTAACAGG
  ACTGATGCCGAAATGTCACCAGGTCCTTTCAGTCTTCACAGTGGAGAACTCTTGG
  30 CCAAAGGTTTTTTGGGGGGAGGAGGAGGAAACCAGCTTTCTGGTTAAGGTTAACA
  - CCAGATGGTTTTTGGGGGGAGGAGGAGGAAACCAGCTTTCTGGTTAAGGTTAACA
    CCAGATGGTGCCCCTCATTGGTGTCCTTTTAAAAAAATATTTACTGTAGTCCAATA
    AGATAGCAGCTGTACAAAATGACTAAAATAGATTGTAGGATCATATGGCGTATA
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- 35 SEQ ID NO: 776

>T97257

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- 40 AATAAGGCAAGTAACTGGGATCCACAATTTATAATACCTGGTCAATTTTTCTGT ATTTAAACCTCTATCATAGGTTTAAGGCCTATTGGGGGGACTTTAATCCCTTACC AAATAAACAGGGGTTTAAAATCACCCTCATGGGGGGCACTGCCCCTTCTGGGGG TTTTCCTTCCTTTGGACTTAAACCAATCTGGGAATGGCTTAGGGGATTTTCCC
- 45 SEQ ID NO: 777

>W96114

GTTACGGAGGCGGCTACGGTGGCCAGAGCAGCATGAGTGGATACGACCAAGTTT TACAGGAAAACTCCAGTGATTTTCAATCAAACATTGCATAGGTAACCAAGGAGC AGTGAACAGCAGCTACTACAGTAGTGGAAGCCCGTGCATCTATGGGCGTGAACG

SEQ ID NO: 778 >AA486836

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**SEQ ID NO: 779** 

20 >L24470 GTGCGCGGAGGGGCCGGCTGGACCACAGCCGGCCCCGATCAGGATCTCC ·CGAGCGGCTCCGTCTTCTGCTCCTCAGAGAGCCCGGCTGGCGGCCTGGGATGACA` AGATGTCTGGACTGCAATCCTGCACAGTTTTGAGAGGGAGATGACTTGAGTGGTT 25 GGCTTTTATCTCCACAACAATGTCCATGAACAATTCCAAACAGCTAGTGTCTCCT GCAGCTGCGCTTCTTCAAACACAACCTGCCAGACGGAAAACCGGCTTTCCGTAT TTTTTCAGTAATCTTCATGACAGTGGGAATCTTGTCAAACAGCCTTGCCATCGCC ATTCTCATGAAGGCATATCAGAGATTTAGACAGAAGTCCAAGGCATCGTTTCTGC TTTTGGCCAGCGGCCTGGTAATCACTGATTTCTTTGGCCATCTCATCAATGGAGCC 30 ATAGCAGTATTTGTATATGCTTCTGATAAAGAATGGATCCGCTTTGACCAATCAA ATGTCCTTTGCAGTATTTTTGGTATCTGCATGGTGTTTTCTGGTCTGTGCCCACTTC TTCTAGGCAGTGTGATGGCCATTGAGCGGTGTATTGGAGTCACAAAACCAATATT TCATTCTACGAAAATTACATCCAAACATGTGAAAATGATGTTAAGTGGTGTGC TTGTTTGCTGTTTTCATAGCTTTGCTGCCCATCCTTGGACATCGAGACTATAAAAT 35 TCAGGCGTCGAGGACCTGGTGTTTCTACAACACAGAAGACATCAAAGACTGGGA AGATAGATTTATCTTCTACTTTTTCTTTTCTGGGGCTCTTAGCCCTTGGTGTTTC ATTGTTGTGCAATGCAATCACAGGAATTACACTTTTAAGAGTTAAATTTAAAAGT CAGCAGCACAGACAAGGCAGATCTCATCATTTGGAAATGGTAATCCAGCTCCTG GCGATAATGTGTCTCCTGTATTTGTTGGAGCCCATTTCTGGTTACAATGGCCAA 40 CATTGGAATAAATGGAAATCATTCTCTGGAAACCTGTGAAACAACACTTTTTGCT CTCCGAATGGCAACATGGAATCAAATCTTAGATCCTTGGGTATATATTCTTCTAC GAAAGGCTGTCCTTAAGAATCTCTATAAGCTTGCCAGTCAATGCTGTGGAGTGCA TGTCATCAGCTTACATATTTGGGAGCTTAGTTCCATTAAAAATTCCTTAAAGGTTG 45 GACAGTAAATCTGTGTGGGGCTAGAACAAAAATTAAGACATGTTTGGCAATATTT CAGTTAGTTAAATACCTGTAGCCTAACTGGAAAATTCAGGCTTCATCATGTAGTT TGAAGATACTATTGTCAGATTCAGGTTTTGAAATTTGTCAAATAAACAGGATAAC TGTACATTTCAACTTGTTTTTGCCAATGGGAGGTAGACACAATAAAATAATGCC 

GAATCATCTGTTGAGGTCTAATGCCTCTACTTGGCCTATTTGCCAGAGAACATCTT AATGCAGCCTGCATAGTGAAATGGTTATTTTGAGATCACCGCTCTGTAGCTAACC CTTATAAACTAGGCTCAGTAAAATAAAGCACTCTTATTTTTTGATCTGGCCTATTT TGCCCCTCATTGTGTAGCCTCAATTAACACATGCATGGTCATGACACCCAGAATT 5 CATGATGGTTTGTTATAACAACCTCTGCATATTCCAGGTCTGGCAGACAGGTTGC CTGACCCTGCAATCCTATCTAGAATGGGCCCATTCTTGTCACATTTGACAAATAG GACTGCCTACATTTATTATTATGAAGGTCGATTGTTGGTAGGAAGTGTTTTTTCATG TGCACAATAATTTTTTAGAGAAACAAAGGCTCTTTCTCAGCACATTGATGGG 10 AGCTACATGCCAATGATAGGTGCAAAGAATATTGGCAAAAGGTGCTTTACCTTG AGCCATTATTTGTGTCAGAGAACAAAAGAAACAGAATCAATATAAATTCAAA GACTATCTGCAGCTAGTGTTTCTTCTTTACACACACATATACACACAGACATCAG AAAATTCTGTTGAGAGCAGGTTCATTAAATTTGTAAGATGGCATATTCTAAAGCC 15 TGTGCTACCAGTACTAAGAGGGGAAGACTGGCAATTTGCCAAGCACTTGGGGAT TATTATAACAATTAACTAGGAGATCAAGAGATAATAATCTCTCCCCAAATTTTCC AATAATAATTGAG

**SEO ID NO: 780** 

20 >T61078

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**SEQ ID NO: 781** >S40706

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30 AGAGACTTAAGTCTAAGGCACTGAGCGTATCATGTTAAAGATGAGCGGGTGGCA GCGACAGAGCCAAAATCAGAGCTGGAACCTGAGGAGAGAGTGTTCAAGAAGGA AGTGTATCTTCATACATCACCACACCTGAAAGCAGATGTGCTTTTCCAGACTGAT  ${\tt CCAACTGCAGAGATGGCAGCTGAGTCATTGCCTTTCTCTTCGGACACTGTCAGCT}$ GGGAGCTGGAAGCCTGGTATGAGGACCTGCAAGAGGTCCTGTCTTCAGATGAAA 35 ATGGGGGTACCTATGTTTCACCTCCTGGAAATGAAGAGGAAGAATCAAAAATCT AGCAGAGGTCACAAGCACCTCCCAGAGCCCTCACTCTCCAGATTCCAGTCAGAG CTCCCTGGCTCAGGAGGAAGAGGAGGAAGACCAAGGGAGAACCAGGAAACGGA AACAGAGTGGTCATTCCCCAGCCCGGGCTGGAAAGCAGCGCATGAAGGAGAAAG 40 AACAGGAGAATGAAAGGAAAGTGGCACAGCTAGCTGAAGAGAATGAACGGCTC AAGCAGGAAATCGAGCGCCTGACCAGGGAAGTAGAGGCGACTCGCCGAGCTCTG ATTGACCGAATGGTGAATCTGCACCAAGCATGAACAATTGGGAGCATCAGTCCC CCACTTGGGCCACACTACCCACCTTTCCCAGAAGTGGCTACTGACTACCCTCTCA CTAGTGCCAATGATGTGACCCTCAATCCCACATACGCAGGGGGAAGGCTTGGAG 45 TAGACAAAAGGAAAGGTCTCAGCTTGTATATAGAGATTGTACATTTATTATTAC

**SEQ ID NO: 782** 

>H25907

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**SEQ ID NO: 783** 

>N90246

- 15 GCTCCCTCCCATGATCATCCTTTTACTTCTACCCCCACCTCCCTTTTAAAACAAAG AGGTTTTCTGGGGTGCAGAGAGGGTTCTCCTGAAAGTGGGCAGAAGCAGCACCG ATAGCCTAGTCTGGCAGGTTGTGGGAAGGGGCGCAGGGAGTGACCATGAGCGAC CTTGGCCCCGTCCTTGCTCCTTGCACCCTGATTGGGGCATGGGGTGAGAGGAGGG ATCAGTCCTTGAATCCCTGAATACTGCAAA
- 20 GAATGCGCTTCTGGTGCCCGGGCAGTGT

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SEQ ID NO: 785 >AA477082

AGAGAGTATATTGCTGCTTTTCTCAGTCACTTTGGCACAGGTGTCGTGGAATATG

ATGCAGAAGGCTTTACAAAACTCACTCTGCTGCTGATGTGGAAAGATTTTTGTTT

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**SEQ ID NO: 786** 

>Z73903

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TTTATGTTTAAGAGGGGCAGTTATAAATGGACACATTGCCCAGAATNTTTTGTAA NATGAAGACCAGCAAATGTAGGCTGATCTCCTTCACAGGATACACTTGAAATAT AGAAGTTATGTTTTAAATATCTCTGTTTTAGGAGTTCACATATAGTTCAGCATTTA TTTTGTTATAATCTTAAGCAACAAAGAAAAACCCTAATATTTGAATCTATTTAT GTCTTTCAATTTAAATTCACTTCAGTTTTTGTTATTGTAATATATTTACTTTTACAT TTCCATGTATTGATGTAGTTAGTCCACATTTAAATTTTTATAGAATTATATAGTTT TTGAAAAATACAGTCAGTAGATGTTTTATTTTTTTAGCTATTCAGTTATGTTTATAA CTCTTTAATGCAATGATTTGTTTTATATTTTGGACTAAGGTTCTTGAGCTTATCTCC CAAGGTACTTCCATAATTTAACACAGCTTCTATAAAAGTGACTTCATGCTTACTT GTGGATCATTCTTGCTGCTTAAGATGAAAAGCATTGGTTTTTTAAAATTAGAGAA TAAAATATGTATTTAAATTTTTGGTGTGTTCACATAAAGNGATGTAGCTAAAATG TTTTCATAGGCTATTATATATNCTCGCAGCATTTCCAGTTAAGAGGATATTAGGT ATATAATTCTCTTCAACCGAATGTCAGATGGTCTTACGCCACAGGGTGCAGGT AACCCTTGGCCTGTAAGCACCACCGATCCAGGGATCATTGTCTAAATAGGTTACT ATTGTTTGTTTCATCTGGAATTCCGAGGGTGCACAAATTACCAGCAGCCTGAGGG AGGTCTTATTGGAAGGCTTCCTGCAGGAGGTGGCATCTGATCCGGGCCTAGACAT GAACAAACGTATGGTGTGGGACCAAGCCTCACATGGTGCATAAATGATCAGAGG CAGGGAAGCTCAGTGAGTGTCTGCTACATGTGAGGCTTCATTCTGAGCACCAACT CTGCGCGCTAATGCCACGAAGTGGAGTCATGATTCTTCCATTTTCCAGGTGAAGA AACCAAAGCACGGGGAGAATTAGCTTGTCCAAGGGCTCATGGCCAGTCGGTGGT GCCACTGGGATTTAACCAGGGATTCCAGCTGTGGAAGCTGCATGCTGCCAGCTGA GCACTCCGTCTGTCCACCCTGGCAAATGACATGCTGAGGGCTGGGGGCAGTCATC TTTCCAGCAGTTGTCGACCAGAAATAATGATAATGCATCAGCCAGGTGGGCAAG AACTCAGACTTTGTGGTCAGACCTCGCTGTGGCAGGGAGCTTAGAAAAGTCACTT TTCCTCCACAGCTGTGTAACATGGGGAAAGGGGGATGCTCAGCTCACCTCATG GCCCTTGTGTGGGGATTTCAAGGGTAAATGCTGGAAAAGCACTGAATATACAGC ACTTAGTAGAGGCTTTGCAAGCGGCTGTCCTCACCCAAACGGATCCTTCCCCAGG AAAGCTCTGCAAATGGAGCCACTTGGCTTTTTGCCTGATGCCATATTCCGTTGTTT AAAACGGCGATCCCAAACTTAAGAGTATGAAAGGGGTGTTTTGCAGACAATGCA TAATTAACCACTTCTAAACACAGGGTGTGCCAGGCTCCCTCAAAGCTGTTAATTA TTCTTCATTAGTGATCTTAAGTTAATGTAAAATAATCCTGGAGCTCTGCCAGAGG TCTTCACGGGCGCCTCTGCGTTCTGCCTTGGTTGTGGGGAGATGCCTCTATAGA CTCTTTAGCTCTGCCCAAAGGCCATTGTGGGAGGCGGCAAGCCCTCTCCCAAGTT CCCAGAGCCCAGCCTCAGGGTACTTTGGTTAATGAAATAGTTCAGGTGGG AAGAAAATATGGAGGAGAGTAGTATACCCATCGCCCAGCTTCCAAAATAAGCTG TTAATGGTGTGGGCCCCTGTGGGCCACCCCACAAATCTTATCCCTTTCCT GCTTCCCCAGAGCTAAGCCTAAACAGTCCCTCTTGTTTGGTGTGTAGATTTCCCAT GCACATTTTTTAAATTGCTGTAGGTACATAGTAAGTGTATATATTTATGGGGTAC GTGAGATGTTTTGACACAGGCACGCAATGTGAAATACACACTTCATGGAGAATTC

SEQ ID NO: 787 >M81882

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ATAATTTTACAAAAGGAAAATATATATATTAAAAAAGATATCCTATTTTGTAACA

SEQ ID NO: 788 >AA401448

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**SEQ ID NO: 789** 

>T84762

ATTCGGCACAGGATGAGCAGAAGAGGTTATCAGCATTAAATTGTTTTGGTTCTAA ATTTGGAACAGTATATATAATTAAAAGTAAGGAACATTAGAGGATTTAATTAGA

- 15 ATAAATACATGTTTTGGAAATACAGTGACCTCTTGCAGTGTCACAAAAGTGCAAA GTGATATTAGCTGTCATCTGCAATACAGAATCTCATTGCTTTTTGCACATGGAGCA TATAGGGAAACTCCANACAGATCACAATGAGGGTTTCTAAATCTGTTGGGGTTCT GTCTTCTATTGGGGTTCTGTGAAGGCAAACCACTGTAGGCTTTAGCTGGGGTTCN GTCCTATGGACTCGTTGGGGGGNATGCCNTG
- 20 GGTTTTTCCATNCTTACCTGGCAGTCTTGGGGGGGGT

GTGAAACTGTTACCCATAAAGTGTAGC

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- - 30 AGTNGGTGCCAGGGTGCAAGTTAGGCTAAAGAAGCCACCACTTATTCCTCTCT TGCCCATTTNTGGGGGGGCAAAGGCCATTTGGTCACCCAAGAGTCTTTCCAGGGG GACCCACAGATATTGCCATGTCCCTNCACACGTCTTTGGNGTCCTTAACN

SEQ ID NO: 791

- 35 >AA424743

- 45 SEQ ID NO: 792

>AA489331

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CAGCAACATGTTTCTCAGCCGAATTACATTCATAGAAAGTGTCCAGATTCTAAAA TCAAATTAAAAGCATGTAATCCAAAGCCTGAAAAAGCAAACAGCTTTAGGGGCT GACTCCATTAGCGTTCCATAGACTGTGCTTTTAACCGTTCAGTTCATGTTTAATGG CCCATCGGTTCCTTACATACCATCAGCTTATGCTGTGGCCAAAAGAAGTGTTCTT GTGGCTTGGTACTCGTCCCTTCAAACAGTAAACAAGAAAGTGCAGACAGTGCTG CCAGAGACAG

**SEQ ID NO: 793** 

>T67104

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TTCTGGTGGCTCAGGAACGTCTTTGGCAACTACAACAGCTGATATTTCAACAGAG
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15 GAACACCTCCAGGCTTCCTCTTTGATGCCACCCACTGGACCTGCCTTGGGGGTCT
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**SEQ ID NO: 794** 

20 >R65792

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AAGAGAGGTCTTGCAGTCTTCAGATGAAACTAAATCTCTAGAAGAGGCACAAGAAT

- 25 GGCTAAAGCAATTCATCCAAGGGCCACCGGAAGTAATTAGAGCTTTGAAAAAAT CTGTTTGTTCAGGCAGAGGCTATATTTGGGGGGAAGCATTACAGAACGAAAGA GATCTTTAGGGNACAGTTTTGGGGTNGGCCNGCAAATTTTAGAGGCTATTTNCT AAGGAAGGGNATTTTATTAATATTTTGGTTTTTCCCG
  - 30 SEQ ID NO: 795

>T90621

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- TATGCCTGACACGCCGGAGGGCTNGAGGGGGAACACACTGAAAGCAGTACCAGG GAGCAGTGCATCTCACAGANCCATTTNTTCATGCCATGAAGTAAACGGTACTTAT ACAAGTGTACAGTGACGTTCCACGNTCCCCATCTAACACGGNTTGCTGGAANTTT ACAGGCAGACTGACGTTTTCTTTCACATGTACTCCAAGTAAATCTGGTTAGTGAT GACCNGGGGGCAGGCGCTGAAGCTTTTCAAAAGCCTTACTTCTTTTATCAGCAGCC
- 40 CGGNTTTTTAT

**SEQ ID NO: 796** 

>AA464067

**SEQ ID NO: 797** >AA291163

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**SEQ ID NO: 798** 

>N53024

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TTTCNTTGTGCATTGCTGACTGTTCCAACATNACAAGTATTATTAAAATTAAATAT - 37x 1 

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25 >AA398230

> GCGGAATGGGAAGCAGTTTATGGAGTTAAGTGGGGCTCTGCTATTTCCCCCAAGA AGGACTCGGAAGATGTTGATTCCAGGGCAGAGTGAGGGCAGACGGGATGAGG  ${\sf CTCTTCTGTAAAGTCCAACAGACGCTCACAGATGCTGGGGAGGCTGGGGACTGCC}$

4: 831

三烯四基式试图2016年代/BCB、1800年代4000年代中代中华的特殊的对象

30 ACCCAGCACACAGGGGCCTCTCCTCACGCTCCCAGGCCACCAGGATGGCCCCC AGGTTCACACACAGGCACACGCACACGCTGCACTCACCACGCACTGAAGGGC ATCACAGCCCCAAGTCTGGGTAAGAAATTCTCCAC

**SEO ID NO: 800** 

35 >H21107

> GTTGCATCATTATACATCACACTGAGTAAGAATCGTTTAGCCATCTACATTCAAT GTTACTGGGTAATATTNNCTCAAATTATAATTCCCAACACTGATTTTACCTGTGA CAAAAGGAACAATAGTAATTCCTTGAAGAAGACTAACTGGNAAAAACATTTNGC TTTTAGTATAAAAGTTCCTAGGNTGCTGAAAGGNACACATACACACCTAAAGAG

TCATGGCCTTCTTAAACAGCTTTCTTAATCCTTTCTGGGAAATATCCTTTGGTTCA 40 TTTTTATTGCCCCTCTCTNGGGCAAAACAAAGTATGTTAACGCAGGNATCAGTGA GTTATNTCCTAGGCACTTGTAAGGCAATATCCTTACCAAGAGGGACCATTCAACT TTTGTAATAATCCGTNAAGCG

45 **SEQ ID NO: 801** 

> zd20g08.s1 Soares fetal heart NbHH19W Homo sapiens cDNA clone IMAGE:341246 3' similar to WP:ZK970.2 CE02402 CLPP-LIKE PROTEASE;, mRNA sequence gi|1365390|gb|W58658.1|W58658[1365390] GCGACCGCCGAGCGACAGATCCAGAACGGCCTGGCCTGCAGCGGTGCCTGACGC

GAACGGCANCCCGGGCTCTCCCGCTCATTCCCATCGTGGTGGAGCAGACGGGTC
GCGGCGAGNCGCCTGATGACATCTACTCGCGGCTGCTGCGGGANGCACTCAGTG
TGCGTCATGGGCCCGATCGATGACAGCGTTGCCAGCCTTGTTATCGCACAGCTCC
TCTTCCTGCAATCCNGAGAGCAACAAGAAGCCCATCCACATGTACATCAACAGC
5 CCTGGTGGTGTGGTGACCGCGGGCCTGGCATCTACGACACGATGCAGTACATCCT
CAACCCGATCTGCACCTGGTGCGTTGGGCCAGGCCGCCAGCATGGGCTCCCTTGC
TTCTCGCCGCCGGAACCCCAGGCATGCGCCACTCGCTCCCCAACTCCGTATCAT
GATCCACCAGCCTCAGGAGGCGCCCG

- 10 **SEQ ID NO: 802** zw32b03.r1 Soares ovary tumor NbHOT Homo sapiens cDNA clone IMAGE:770957 5', mRNA sequence gi|2112210|gb|AA428170.1|AA428170[2112210] CTCATGCGTGAGACTGGGTGTACAAACTCATCCTCTTTTAATGGCATTTCTCTTTA 15 AACTATGTTCCTAACAAAATGAGATGATAGGATAGATCCTGGTTACCACTCTTTT GCTGTGCACATACGGGCTCTGACTGGTTTTAATAGTCACCTTCATGATTATAGCA ACTAATGTTTGAACAAGCTCAAAGTATGCAATGCTTCATTATTCAAGAATGAAA AATATAATGTTGATAATATATATAAGTGTGCCAAATCAGTTTGACTACTCTCTGT 20 TATTTCTCTATTTCATAATCAGTAATAGTGTCATATAAACTCATTTATCTCCTCTT CATGGCATCTTCAATATGAATCTATAAGTAGTAAATCAGAAAGTAACAATCTATG GCTTATTTCTATGACAAATTCAAGAGCTAGAAAAATA
  - **SEQ ID NO: 803**
- ab35g03.s1 Stratagene HeLa cell s3 937216 Homo sapiens cDNA clone IMAGE:842836 3' similar to gb:M93056 LEUKOCYTE ELASTASE INHIBITOR (HUMAN);, mRNA sequence gi|2216491|gb|AA486275.1|AA486275[2216491]
- 35 CCAGACAGATCAGCCTTGCTACTGTTAAAGAGATCCTGCACACCTAGGCGGCG AGGTCGGAGTTGAGAGTGTAACTCTCTTCCAGTTTGAACCTGGGCAAGCTGACAT TAACTTCAATGAAATCGAGATTCTCAGGTTTAG

## (19) World Intellectual Property Organization International Bureau





(43) International Publication Date 26 September 2002 (26.09.2002)

**PCT** 

# (10) International Publication Number WO 02/074979 A3

(51) International Patent Classification<sup>7</sup>: C12Q 1/68

(21) International Application Number: PCT/US02/08456

**(22) International Filing Date:** 20 March 2002 (20.03.2002)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:

60/276,947 20 March 2001 (20.03.2001) US

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(81) Designated States (national): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZM, ZW.

(84) Designated States (regional): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

#### Published:

with international search report

(88) Date of publication of the international search report: 13 March 2003

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.



(54) Title: EXPRESSION PROFILES AND METHODS OF USE

(57) Abstract: The present invention relates to gene expression profiles, algorithms to generate gene expression profiles, microarrays comprising nucleic acid sequences representing gene expression profiles, methods of using gene expression profiles and microarrays, and business methods directed to the use of gene expression profiles, microarrays, and algorithms. The present invention further relates to protein expression profiles, algorithms to generate protein expression profiles, microarrays comprising protein-capture agents that bind proteins comprising protein expression profiles, methods of using protein expression profiles and microarrays, and business methods directed to the use of protein expression profiles, microarrays, and algorithms.

### INTERNATIONAL SEARCH REPORT

International application No.

PCT/US02/08456

A. CLASSIFICATION OF SUBJECT MATTER  IPC(7) : C12Q 1/68 US CL : 435/6			
According to International Patent Classification (IPC) or to both national classification and IPC			
B. FIELDS SEARCHED			
Minimum documentation searched (classification system followed by classification symbols)			
U.S. : 435/6			
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched			
Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)			
Electronic data base consumed during the international search (hame of data base and, where practicable, search terms used)			
C. DOCUMENTS CONSIDERED TO BE RELEVANT			
Category *	Citation of document, with indication, where a	opropriate of the relevant passages	Relevant to claim No.
A, E	POOLE et al. Altered Patterns of Cellular Gene Exp		1, 3-4, 6-90
71, —	Endothelial Cells Infected with Kaposi's Sarcoma-associated Herpesvius. Journal of		
	Virology. April 2002, Vol 76, No. 7, pages 3395-34		
Α, Ρ	US 6,316,197 B1 (DAS et al) 13 November 2001 (1)	3.11.2001), entire document.	1-90
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71, <b>L</b>	US 0,403,510 B1 (SKALITER et al) 11 June 2002(11.00.2002), Chine document.		1-50
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Further	documents are listed in the continuation of Box C.	See patent family annex.	
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"A" document	defining the general state of the art which is not considered to be	date and not in conflict with the applic principle or theory underlying the inve	ation but cited to understand the
of particular relevance			
"E" earlier application or patent published on or after the international filing date		"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone	
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		,	
"P" document published prior to the international filing date but later than the priority date claimed		"&" document member of the same patent i	amily
Date of the actual completion of the international search Date		Date of mailing of the international search	ch report
		06 NOV 2002	
30 September 2002 (30.09.2002)  Name and mailing address of the ISA/US		Authorized officer	
	nmissioner of Patents and Trademarks		100
Box PCT Washington, D.C. 20231		Shubo "Joe" Zhou Wella (	allenson
		Telephone No. (703)-308-0196	
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